

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number

TO: Minh-Tam Davis

Location: REM-3A24&3C18

Art Unit: 1642

Thursday, March 09, 2006

Case Serial Number: 10/001469

From: Alex Waclawiw

Location: Biotech-Chem Library

Rem 1A71

Phone: 272-2534

Alexandra.waclawiw@uspto.gov

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STIĆ-Biotech/ChemLib

181717

From:

Chan, Christina

Sent:

Wednesday, March 08, 2006 2:25 PM Davis, Minh-Tam; STIC-Biotech/ChemLib

To: Subject:

RE: Rush search request for 10/001469

Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89

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----Original Message----

From:

Davis, Minh-Tam

Sent:

Wednesday, March 08, 2006 11:19 AM

To:

Chan, Christina

Subject:

Rush search request for 10/001469

Please search in commercial database, issued patent files, PGPUB and interference: The protein of SEQ ID NO:2866

Thank you.

MINH TAM DAVIS

ART UNIT 1642, ROOM 3A24, MB 3C18

272-0830

************************** Searcher Phone: Date Searcher Picked up Date completed: Searcher Prep Time:	Technical Info. Specialist
Online Time:	· · ·

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The invention relates to a novel composition comprising: a substance that modulates the status of a protein comprising a sequence with a fully defined 2466 or 3136 amino acid sequence given in the specification; or a molecule that is immobilised by a protein comprising a sequence with a fully defined 2466 or 3136 amino acids, where the status of a cell expressing the protein is modulated. The invention further relates to: an antibody; a non-human transgenic animal or hybridoma that produces the antibody; a method of delivering a cytotoxic or diagnostic agent to a cell that expresses the protein; a polymucleotide that encodes the protein; a method for inhibiting the growth of cancer cells that express the protein; a method for generating a mammalian immune response directed to the protein; a method detecting in a sample the presence of a 101P3A11
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Challita-Eid
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Matches 318;
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New composition comprising 101P3A11-related protein, useful preventing or treating cancer e.g., stomach, cervix, uterus, prostate, colon, kidney or breast cancer.
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31-OCT-2001; 2001US-00001469.
14-DEC-2001; 2001US-00017666.
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Challita-Eid
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Pred. No. 1e-
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                                                                   The invention relates to a novel composition comprising: a substance that comodulates the status of a protein comprising a sequence with a fully confined 2466 or 3136 amino acid sequence given in the specification; or a confidence that is immobilised by a protein comprising a sequence with a confidence of the comprising a sequence with a confidence of the protein is modulated. The invention further relates to: an expressing the protein is modulated. The invention further relates to: an contibody; a method of delivering a cytotoxic or diagnostic agent to a contibody; a method of delivering a cytotoxic or diagnostic agent to a coll that expresses the protein; a polymucleotide that encodes the correctin; a method for inhibiting the growth of cancer cells that express the protein; a method for generating a mammalian immune response directed to the protein; a method detecting in a sample the presence of a 101P3A11 correlated protein or polymucleotide; and a method for monitoring one or corrected of having cancer. The novel composition has cytostatic cactivity. The composition is useful for preventing or treating cancer cells, stomach, cervix, uterus, rectum, prostate, colon, kidney or breast of the invention.
Sequence 318 AA;
                                                       of the invention.
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밁 δ 밁 Ś 밁 5 밁 S 8 片 Ś Query Match Best Local S Matches 318 241 181 181 121 121 318; 13 13 Similarity IRORILRLFHVATHASEP 318 GTCVSHVCAVFIFYVPFIGLSMYHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKS GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKB YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS MAPDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVPIKQLPFCRSNILSHS PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA MMVDPNGNESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE MMYDPNGNESSATYFILIGLEGLEEAQFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHE YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTRBAQAKAF IRQRILRLFHVATHASEP 100.0%; Score 1622; ilarity 100.0%; Pred. No. 1e-Conservative 0; Mismatches 12; DB 7; 7; 0 Indels Length 318; 0 120 300 300 240 180 180 120 60 0

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RESULT 4
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DT 22-P
CX
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DX Unid
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PN W020
PD 21-P
                                                                                               22-APR-2004
                                                                                                                ADI09912;
                                                                                                                             ADI09912 standard; protein; 338
                                                                                               (first
                                                                                               entry)
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transgenic animal; cytotoxic; cancer; immune; 101P3A11; cytostatic; stomach; cervix; uterus; rectum; prostate; colon; kidney; breast.

Optimised 101P3A11v1 gene protein.

15-MAY-2002;

2002WO-US015520

GCRP-5; G protein-coupled receptor protein; antipsoriatic; antirheumatic; immunosuppressive; antiasthmatic; antianentc; antiarteriosclerotic; anti-inflammatory; anti-inflammatory; antigout; thyromimetic; haemostatic; virucide; hepatotropic; osteopathic;

antiparasitic;

immunostimulant.

10-AUG-2000 AAY92365;

(first entry)

receptor protein

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel composition comprising: a substance that CC modulates the status of a protein comprising a sequence with a fully CC defined 2466 or 3136 amino acid sequence given in the specification; or a CC molecule that is immobilised by a protein comprising a sequence with a CC fully defined 2466 or 3136 amino acids, where the status of a cell CC expressing the protein is modulated. The invention further relates to: an CC antibody; a non-human transgenic animal or hybridoma that produces the CC antibody; a method of delivering a cytotoxic or diagnostic agent to a CC cell that expresses the protein; a polynucleotide that encodes the CC to the protein; a method for generating a mammalian immune response directed to the protein; a method detecting in a sample the presence of a 10193A11 CC related protein or polynucleotide; and a method for monitoring one or CC correction or polynucleotide; and a sample from a patient having CC or suspected of having cancer. The novel composition has cytostatic cativity. The composition is useful for preventing or treating cancer ce.g., stomach, cervix, uterus, rectum, prostate, colon, kidney or breast cytoscape. This sequence represents a 10193A11 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 318
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31-OCT-2001;
14-DEC-2001;
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                 AAY92365
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Challita-Eid PM;
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                 standard;
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2001US-00001469.
2001US-00017666.
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                protein; 318
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Pred. No. 1.1e-169;
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06-OCT-1998;
11-MAY-1999;
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Corley
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                                                      Y YT,
                                        2000-328934/28.
DB; AAA09351.
                                                                                INCYTE PHARM INC.
                                                           Yue H, Lal P,
Guegler KJ,
                                                                                             98US-00167219.
98US-0172211P.
99US-0133585P.
                                                                                                                         99WG-US023317.
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57
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272
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note= "G-protein coupled receptor"
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"G-protein_coupled_receptor"
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                                                           Bandman O,
Gorgone GA,
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                                                           Au-Young J
Baughn MR,
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Novel human G-protein coupled receptor proteins used in the diagnosis, treatment and prevention of nervous system disorders, autoimmune/inflammatory disorders, and cell proliferative disorders su

disorders such

as cancer

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Matches 317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorders, complications of cancer, hemadialysis and extracorporeal circulation, and cell proliferative disorders. They are also used to treat or prevent disorders associated with decreased or increased
                       Xu J, Dillon DC, Mitcham Kalos MD, Fanger GR, Day Wang A, Meagher MJ;
WPI; 2001-425873/45
                                                                                                                                16-JAN-2001; 2001WO-US001574.
                                                                                                                                                                                                                                   Human; prostate cancer; prostate-specific; diagnosis; vaccine;
cytostatic; gene therapy; metastasis.
                                                                                                                                                                                                                                                                           P835P amino acid sequence
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                                                                                                                                                                                                                                                                                                                                                          AAM01306 standard;
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                                                                            (CORI-) CORIXA CORP.
                                                                                                     14-JAN-2000; 2000US-00483672
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99.7%;
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Pred. No. 2.2e-169;
1; Mismatches 0;
                                       Harlocker SL, Jiang Y, Retter MW, Stolk JA,
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The present invention describes polynucleotide sequences (I) which encode CC prostate-specific proteins (II). (I) and (II) have cytostatic activity, CC and can be used in vaccine production and gene therapy. (I), (II), CC antibodies to (II), fusion proteins comprising (II), and isolated T cells CC prepared using (I) or (II) are used treat cancer in a patient. (I) and CC cancer that is diagnosed or treated is particularly prostate cancer. The CC and (II) can be used in vaccines. The antibodies or (I) can be used for CC concer that is diagnosed or treated is particularly prostate cancer. (I) CC and (II) can be used in vaccines. The antibodies or (I) can be used for CC woiltowing the progression of cancer in a patient. (I) and (II) can also CC be used to improve diagnostic and therapeutic methods for prostate CC cancer. They can indicate the level of metastasis as well as the prostate CC polynucleotide and amino acid sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; Page 534-535; 543pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotide encoding a prostate-specific protein, monitoring and treating prostate cancer in a patient and
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Sequence 318

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밁 S 맑 S 밁 Ś 문 5 밁 Ś 片 S Query Match Best Local S Matches 317 301 301 241 241 181 181 121 al Similarity 317; Conserv 61 61 IRQRILRLFHVATHASEP 318 GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKE 300 YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120 MMVDPNGNESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE MMVDPNGNESSATYFILIGLPGLEEAQFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHE YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS IRORILRLFHVATHASEP GTCVSHVCAVF1FYVPF1GLSMVHRFSKRRDSPLPV1LAN1YLLVPPVLNP1VYGVKTKB 99.8%; Score 1619; DB 4; ilarity 99.7%; Pred. No. 2.2e-169; Conservative 1; Mismatches 0; 318 Length 318; Indels 0 240 180 180 60 240 0

AAU69951 standard; protein; 318

AAU69951;

30-JAN-2002 (first entry)

Human prostate cDNA encoded protein

Human; prostate cancer; cytostatic; immunostimulant; tumour; immunogen

WO200173032-A2

04-OCT-2001.

27-MAR-2001; 2001WO-US009919

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09-MAY-2000

12-MAY-2000

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27-JUN-2000

09-AUG-2000

09-AUG-2000

06-SEP-2000

02-CCT-2000

02-CCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to isolated prostate-specific polynucleotides, polypeptides, fusion proteins of the polypeptides, antibodies raised against the polypeptides (or antigenic epitopes derived from them) and antigen-presenting cells expressing the polypeptides. The antibodies are useful for detecting the presence of cancer, especially prostate cancer. The polypeptides, polynucleotides and the antigen-presenting cells are useful for stimulating and/or expanding T cells specific for a tumour protein, and for inhibiting the development of cancer especially prostate cancer. Compositions comprising the polynucleotide and/or polypeptide are useful for stimulating an immune response, and for treating cancer. The oligonucleotide is useful for detecting cancer. The present sequence is a prostate specific polypeptide of the invention
 AAG71659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 318 AA;
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N-PSDB; AAS64174, AAS64176, AAS64177.
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Li SX,
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GR, Retter MW, Stolk JA,
Wang A, Skeiky YAW, Hep
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standard; protein; 318
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2000US-00568100.
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2000US-00636215.
2000US-00636215.
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2000US-00679426.
2000US-00685166.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 1619; DB 4;
Pred. No. 2.2e-169;
1; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 318;
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Best Local Sim
Matches 317;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is an olfactory receptor which is encoded by one of a number of novel polynucleotides. The polynucleotides can be used in screening for olfactory agonists and antagonists. The methods allow for the determination of primary scents and the identification of the odour receptors used to detect these primary scents. The methods also enable determination of secondary scents and the identification of combinations of odour receptors that are involved in detecting such secondary scents. This enables the construction of a scent representation (also called a scent fingerprint or scent profile), which may be used to re-create and edit scents. Libraries of olfactory receptors are useful for determining the interaction pattern of a composition with the receptors, and can be used to determining differences in the olfactory faculties of different and of the control of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 11; Page 826-827; 1857pp; English.
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24-FEB-2000;
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Pred. No. 2.2e-169;
1; Mismatches 0;
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99US-00428114.
99US-00429517.
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274. .295
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/note= "Third BC loop"
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/note= "Second EC loop"
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/note= "First IC loop"
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"Second IC loop"
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to human G protein-coupled receptor (GPCR) and their corresponding DNA molecules. GPCR is also referred as seven transmembrane receptor. G protein-coupled receptor protein is useful for treating neurological disorder, particularly schizophrenia. GPCR protein is also useful for identifying compounds useful for treating schizophrenia. These compounds are also useful for treating other neurological and psychiatric diseases, e.g. depression, anxiety, bipolar disease, affective disorders, attention deficit hyperactivity disorder/attention deficit disorder, epilepsy, neuritis, neurostathe and senile dementia. The invention also provides genetic screening procedures that entail analysing a person's genome with respect to GPCR. The vectors are useful for the recombinant production of the GPCR's. The present sequence is human CON198 G protein-coupled receptor (GPCR) protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seven transmembrane receptor polypeptides and polynucleotides, useful iteating neurological or psychiatric disorders, e.g. schizophrenia, as well as for identifying compounds useful for treating schizophrenia.
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N-PSDB; AAD06504.
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12-JAN-2000; 2000US-00481794.
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Pred. No. 2.2e-169;
1; Mismatches 0;
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RESULT 10
ABB44533
ID ABB44
XXXXXXXX
                             28-JAN-2002
                                             ABB44533;
                                                          ABB44533 standard;
             Human GPCR7a polypeptide SEQ ID NO
                             (first entry)
                                                          protein;
               26
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Human; GCPR; G-coupled protein-receptor; cardiant; antiarteriosclerotic;

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The invention relates to nucleic acid sequences (ABA81529-ABA81522) that carcode G-coupled protein-receptor related polypeptides (ABB44522-CC ABB44543). The isolated polypeptide having a sequence differing by no CC more than 15 % of amino acid residues from one of 22 amino acid sequences (or mature forms of the sequences), fully defined in the specification CC and corresponding to human G-protein coupled receptor X (GPCRX) CC polypeptides. The polypeptides have potential cardiant, CC antiarteriosclerotic, anabolic, cytostatic and antiviral activity. The CC polypeptides can be administered therapeutically, especially using gene CC therapy and expressing the encoding DNA in vivo, to treat or prevent CC GPCRX-associated disorders, especially in humans. For example, they can be used to treat/prevent cardiomyopathy, atherosclerosis, disorders (crelated to signal processing and metabolic pathway modulation (e.g. cobssity, anorexia), diabetes, osteoporosis, Crohn's disease, multiple CC disease, Parkinson's disorder, Huntington's disease, immune disorders, haematopoletic disorders, developmental diseases, neurological disorders, bacterial, fungal, protozoal and viral infections (e.g. with human communed file of the polypeptide in mammals (especially humans) by to determine the presence of or predisposition to a disease associated with altered levels of the polypeptide in mammals (especially humans) by catecting alterations in polypeptide expression levels relative to control samples. They are useful to identify agents binding polypeptide colors, cellular receptors or downstream effectors) and/or agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-MAR-2000;
05-APR-2000;
06-APR-2000;
06-APR-2000;
06-APR-2000;
06-APR-2000;
06-APR-2000;
06-APR-2000;
06-APR-2000;
11-AUG-2000;
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11-AUG-2000;
11-AUG-2000;
11-AUG-2001;
23-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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Padigaru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New human G-protein coupled receptor X, GPCRX, polypeptide useful in treatment or prevention of GPCRX associated disorders e.g. cardiomyopathy or atherosclerosis, and to screen for antagonists and agonists useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapeutically.
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DB; ABA81542.
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2000US-0195068P.
2000US-0195069P.
2000US-0195070P.
2000US-0195150P.
2000US-0219855P.
2000US-021985P.
2000US-0221284P.
2000US-0221384P.
2000US-0224584P.
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2000US-0194614P.
2000US-0195063P.
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Mishnu VS,
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Tchernev VT,
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Spytek KA,
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Li L, Baumgartner
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RESULT 11
ABUT1842
ID 71842
ID 7847
AC ABUT1
XX ABUT1
XX Prost
XW Prost
XW Prost
XW Prost
XW PSWA
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Best Local
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04-OCT-2000; 2000US-00679272
28-MAR-2001; 2001US-00822827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prostate cancer; vaccine; gene therapy; cytostatic; fusion protein; immunogen; cancer; prostate specific antigen; PSA; prostatic acid phosphatase; PAP; prostate specific membrane antigen
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(MITC/)
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317; Conserv
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FANGER G R.
RETTER M W.
STOLK J A.
DAY C H.
VEDVICK T S.
CARTER D.
                                                                                                                                     MITCHAM J L.
HARLOCKER S
JIANG Y.
KALOS M D.
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DILLON D
MITCHAM J
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nilarity 99.7%;
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Pred. No. 2.2e
1; Mismatches
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RESULT 1:
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Best Local S
Matches 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention describes a fusion protein comprising at least one amino acid sequence of immunogenic portions of any of the 3 sequences not defined in the specification, or sequences having at least 70 or 90 % sequence identity to any one of the 35 sequences defined in the USPTO we site, which is encoded by any of the 4 nucleotide sequences not defined in the specification. The fusion protein, composition and methods are useful for diagnosing, preventing and/or treating cancer, particularly prostate cancer. The proteins are useful as markers to indicate the presence or absence of cancer. This is the amino acid sequence of a prostate cancer therapy associated protein. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the US patent office at analyze the protein.
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Li SX, Wan
  AAU24561;
                           AAU24561
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(SKEI/)
(HEPL/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MCNEILL P D.
HOUGHTON R L.
Y DE BASSOLS C
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HEPLER W T.
HENDERSON R A.
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SKEIKY
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                           standard; protein; 318
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     99.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 1619; Di
Pred. No. 2.2e
1; Mismatches
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Carter D;
Hural J;
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Best Local Similarity
Matches 317; Conserv
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24-MAR-2000;
12-APR-2000;
12-APR-2000;
24-APR-2000;
26-MAY-2000;
23-JUN-2000;
                                                                                                                                                                                                                                                                                                               The invention relates to nucleic acids encoding human olfactory receptors, OR, (a G protein-coupled receptor, GPCR). The OR's specifically recognise molecules, odourants, that elicit specific olfactory sensation. The human olfactory receptors and polynucleotides encoding them are useful for screening a library of chemical compounds for compounds that are involved in olfactory sensation. Modulators of their activity are useful for pharmacological and genetic modulation of olfactory signalling pathways. Therefore, they can be used in the food, pharmaceutical and cosmetic industries to customise odours and fragrances. The present sequence is a human olfactory receptor of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acids encoding human olfactory G protein-coupled receptors, useful for screening for compounds involved in olfactory sensation, the compounds can be used in the food, pharmaceutical and cosmetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; olfactory receptor; G protein-coupled receptor; GPCR; odourant; food additive; cosmetic; fragrance; pharmaceutical additive.
                                                                                                                                                                                                                                                                           Sequence 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
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MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS
                                                                                                                           PMYIFICMLSGIDILISTSSMPKMLAIFWFWISTTIQFDACLLQIFAIHSLSGMBSTVLLA
                                                                                                                                                                                MMVDPNGNESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE
                                                                                                                                                                MMVDPNGNESSATYFILIGLPGLEBAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE
                                                      MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS
                                                                                                           PMY1FLCMLSGIDILISTSSMPKMLA1FWFNSTT1QFDACLLQMFA1HSLSGMESTVLLA
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2000US-0192033P.

2000US-0198474P.

2000US-019933SP.

2000US-0207702P.

2000US-0213849P.

2000US-0230732P.

2000US-0230732P.

2000US-0236534P.

2000US-0236534P.

2000US-0236534P.

2000US-0266862P.
                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                           AA;
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RESULT 13
ABB95411
ID ABB95
XX ABB95
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01-AUG-1997; 97US-00904804.
10-FEB-1998; 98US-00020956.
25-FEB-1998; 98US-00159612.
14-JUL-1998; 98US-00159812.
15-JAN-1999; 99US-00232149.
09-APR-1999; 99US-00232149.
13-JUL-1999; 99US-00232616.
13-JUL-1999; 99US-00443686.
13-JAN-2000; 2000US-00433313.
18-NOV-1999; 99US-004436857.
27-MAR-2000; 2000US-00536857.
19-MAY-2000; 2000US-00568100.
12-MAY-2000; 2000US-0056810.
12-MAY-2000; 2000US-00579737.
13-JUN-2000; 2000US-00569793.
10-AUG-2000; 2000US-00565729.
06-SEP-2000; 2000US-0065783.
10-OCT-2000; 2000US-0065783.
10-OCT-2000; 2000US-0065783.
                                                                                                 (XUJJ/)
(DILL/)
(MITC/)
(HARL/)
(JIAN/)
(KALO/)
(FANG/)
(RETT/)
(STOL/)
(DAYC/)
(CART/)
(LISX/)
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(SKEI/)
                                  (HEND/)
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                                                ) CARTER D.
) LI S X.
) WANG A.
) SKEIKY Y A W.
) HEPLER W T.
                                                                                                                                FANGER G R.
RETTER M W.
STOLK J A.
DAY C H.
VEDVICK T S.
 Dillon DC,
                                                                                                                                                                                                                    JIANG Y.
KALOS M D.
                                                                                                                                                                                                                                                                     MITCHAM J L
                                                                                                                                                                                                                                              MITCHAM J L.
HARLOCKER S L.
                                 HENDERSON R A.
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Mitcham JL,
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RESULT 14
ABP95674
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Best Local Similarity 99.7%;
Matches 317; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides prostate-specific coding sequences and their encoded proteins. These can be used in the diagnosis and treatme of cancers, particularly prostate cancer. The present sequence is a protein described in the invention
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Li SX, Wang A, Skeiky YAW, Hepler WT, Henderson
                                                                                                                                                                                              brap
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                            04-AUG-2000; 2000JP-00237818.
13-FEB-2001; 2001JP-00034434.
                                                                        30-JUL-2001; 2001WO-IB001446
                                                                                                       28-FEB-2002.
                                                                                                                                                                   Homo
                                                                                                                                                                                                               Human;
                                                                                                                                                                                                                                          Human GPCR polypeptide SEQ ID NO 158
                                                                                                                                                                                                                                                                          06-MAR-2003
                                                                                                                                                                                                                                                                                                        ABP95674;
                                                                                                                                                                                                                                                                                                                                   ABP95674 standard; protein;
(NISC-) JAPAN SCI & TECHNOLOGY CORP.
                                                                                                                                     WO200216548-A2
                                                                                                                                                                   sapiens.
                                                                                                                                                                                            n; GPCR; G protein coupled receptor; signal transduction; olfactory;
development; gustatory; taste; fragrance; receptor.
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Pred. No. 2.2e-169;
1; Mismatches 0;
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enderson RA;
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AAU95746
ID AAU95746
AC AAU95
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Best Local
                                                                                 Human; olfactory and pheromone G protein coupled; receptor; GPCR; tranquilizer; antidepressant; neuroleptic; endocrine; anabolic; anorectic; taste; fragrance; food additive; cosmetic; cell migrat sterility; psychotic disorder; neurological disorder; anxiety; schizophrenia; manic depression; depression; axonal growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABP95942) by extracting open-reading frames containing 6-8 transmembrane domains with 250-1000 amino acid residues to give a gene homologous with a known GPCR gene. The receptor proteins and encoded genes are useful for studying in vivo signal transduction mechanism and identifying targets for drug development e.g. based on olfactory and gustatory receptors in form of agonists and antagonists by screening intrinsic and extrinsic ligands as bitter taste inhibitors, taste enhancers and fragrance improvers. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                              Human olfactory and pheromone G protein-coupled receptor #233.
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU95746 standard; protein;
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                                                      menstrual cycle; appetite
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ABP95942)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                          (first
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                                                            sexual motivation;
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Pred. No. 2.
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                                                            sexual attraction;
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                                                                                                                                              cell migration;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to olfactory and Pheromone G-protein coupled CC receptor (GPCR) or a protein 95% identical to the GPCR, a specific active CC portion and its encoding polymuclectide. Also included are an agonist, CC antagonist or inhibitor of the GPCR or the polymuclectide, a vector CC comprising the polymucleotide, a cell transformed by the vector, a non-clouding the receptor and screening (detection and possibly, recovering) CC encoding the receptor and screening (detection and possibly, recovering) CC inhibitors of natural compounds to the GPCR. The receptor-derived CC agonists, antagonists, inhibitors or compounds are used as an CC improvement, elimination or substitution of an existing taste and/or a fragrance of (or in) the food and/or cosmetic products. They can also be CC used in the preparation of medicament in the treatment and/or prevention of amamalian disorder, such as cell migration, sterility, psychotic and neurological disorders, including anxiety, schizophrenia, manic CC depression, depression, for promoting axonal growth, nerve cell CC endocrine functions, hormone production and the menstrual cycle, for the prevention or the treatment by stimulation of appetite, sexual attraction, segments of appetite, sexual communication between organisms. The present cc sequence is a human olfactory and pheromone GPCR protein sequence
                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 317; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel pheromone G-protein coupled receptor and receptor-derived agoniant antagonists or inhibitors useful in food or cosmetic products or in the treatment or prevention of neurological disorders such as anxiety and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-330013/36.
N-PSDB; ABK68633.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 710-711; 833pp; English.
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GTCVSHVCAVFIFYVPFIGLSMYHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKB
                                                                    YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTRBAQAKAF
                                                                                           YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF
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                                                                                                                                             MAFDRYVAI CHPLRHATYLTLPRVTKI GVAAVVRGAALMAPLPVFI KQLPFCRSNI LSHS
                                                                                                                                                                                                                     PMY1FLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQMFAIHSLSGMESTVLLA
                                                                                                                                                                                                                                                                                            MMVDPNGNESSATYFILIGLPGLEBAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE
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                                                                                                                                                                                                                                                                                                                                                                                99.8%;
                                                                                                                                                                                                                                                                                                                                                             Score 1619; DB 5;
Pred. No. 2.2e-169;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                     Length 318;
                                                                                                                                                                                                                                                                                                                                                                0,
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                                                                                                                                                          Query Match
Best Local 9
                                                                                                                                            Matches
                                                                                                                                                                                                                                         The invention relates to a method of representing sensory perception of one or more odourants. The method comprises: (a) providing a representative class of n olfactory receptors or ligand binding domains (LBD) of these receptors; (b) measuring values X1 to Xn representative of at least one activity of one or more odourants selected from: (i) binding one or more odourants to the LBD of at least one of the n olfactory receptors; (ii) activating at least one of the n olfactory receptors with the one or more odourants; and (iii) blocking at least one of the n olfactory receptors with the one or more odourants; and (c) generating a representation of the sensory perception from the values X1 to Xn. The design and formulation of odour and taste compositions. AAU85140-AAU85393 represent human olfactory G-coupled receptor amino acid sequences of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Representing sensory perception of one or more odorants identification and design of tastes and odors comprises representative group of n olfactory receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G-coupled olfactory receptor
                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 77; 182pp; English.
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N-PSDB; ABK37540.
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13-MAR-2001; 2001US-00804291.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; olfactory G-coupled receptor; sensory perception of odourant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SENO-) SENOMYX INC.
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  121
                                                                                                                                           al Similarity
317; Conserv
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                                                                                                     MMVDPNGNESSATYFILIGLEGLEBAQFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHE
                                            PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
                                                                                                                                                                                                     318 AA;
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  MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS 180
                                                                                      MMVDPNGNESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE
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                                                                                                                                              Conservative
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99.7%;
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                                                                                                                                                          Score 1619; DB 5;
Pred. No. 2.2e-169;
                                                                                                                                            Mismatches
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                                                                                                                                                                         Length 318;
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Claim 11; Fig

2; 74pp;

English

disorders.

The invention relates to a new isolated G protein-coupled receptor

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RESULT 17
ABO19491
ID ABO19491
AC ABO19
AC CODIN
ACCOMMAN ABO19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chagas's disease; post-inflammatory pseudotumour; infection; asthma; benign prostatic hyperplasia; cancer; anorexia; Parkinson's disease; acute heart failure; hypotension; hypertension; osteoporosis; ulcer; neurological disorder; AlDS; allergy; anaemia; atherosclerosis; crohn's disease; atopic dermatitis; diabetes mellitus; emphysema; crave's disease; lupus erythematosus; multiple sclerosis; vaccine; myasthenia gravis; osteoarthritis; osteoporosis; pancreatitis; rheumatoid arthritis; psoriasis; Sjogren's syndrome; trauma; autoimmune thyroiditis; Alzheimer's disease; amnesia; bipolar disorder; autoimmune thyroiditis; Alzheimer's disease; amnesia; bipolar disorder;
                                                                                                                                           New human G-protein coupled receptor polypeptide, HGPRBMY4, expressed highly in prostate, colon and lung, useful for treating, preventing cancer, immune, neurological, cardiovascular, colon and lung-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-SEP-2000; 2000US-0235839P-
16-JAN-2001; 2001US-0261776P-
13-JUL-2001; 2001US-0305351P-
17-AUG-2001; 2001US-0313202P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amyotrophic lateral sclerosis; schizophrenia; Tourette's disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; receptor; G protein-coupled receptor;
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                                                                                                                                                                                                                                                                                                                                    Barber
                                                                                                                                                                                                                                                                                                                                                                 Feder JN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (FEDE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HAWK/)
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DB; ACD29000.
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) MINTIER G.
) RAMANATHAN (
) HAWKEN D R.
) CACACE A.
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Kornacker MG;
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RESULT 18
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Best Local Similarity
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Cytostatic; gene therapy; prostate-specific protein; PSP; human;
                                                           Prostate tumour specific protein SEQ ID 920.
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Pred. No. 2.2e-169;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to novel prostate-specific proteins (PSP) and their coding sequences. The PSPs and their coding sequences are useful for stimulating an immune response in a patient, or for treating prostate cancer in a patient and for determining, detecting or diagnosing the presence of a cancer in a patient. The present sequence was used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New prostate-specific proteins and genes, useful in gene therapy, particularly for stimulating an immune response in a patient, or prostate cancer in a patient, as well as for diagnosing prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xu J, Dillon DC, Mitcham JL, Harlocker SL, Jiang Y, Henderson
KaloB MD, Fanger GR, Retter MM, Stolk JA, Day CH, Vedvick TS;
Carter D, Li SX, Wang A, Skeiky YAW, Hepler WT, Hural J;
Mcneill PD, Houghton RL, Vinals Y De BassolsC, Foy TM, Watanabe
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29-JUN-2001; 2001US-00895814.
10-DEC-2001; 2001US-00012896.
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 318
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                IRQRILRLFHVATHASEP
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Pred. No. 2.2e-169;
"" amatches 0;
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240 240 180 180 120 120 60

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CC any one of 1601 sequences (see ABP82019 to ABP83319) of 12-24 amino CC any one of 1601 sequences (see ABP82019 to ABP83319) of 12-24 amino CC acids. Also described: (1) an assay for the detection of a particular GC protein-coupled receptor (GPCR) or a candidate polypeptide in a sample; CC and (2) an isolated antibody having high specificity and high affinity or CC avidity for a particular GPCR. (1) can be used as GPCR modulators and in CC gene therapy. The antigonic peptides for GPCRs are useful in detecting an CC antibody against a particular GPCR, and in the production of specific CC antibodies. The peptides and antibodies are also useful for detecting the presence or absence of corresponding GPCRs. The antigonic peptides for CFCRs and antibodies are useful for disgnosing and designing drugs for treating immune-related disease, growth-related diseases, cell CC regeneration-related disease, e.g. AIDS, Alzheimer's disease, CC disease, or autoimmune diseases, e.g. AIDS, Alzheimer's disease, CC atherosclerosis, bacterial, fungal, protozoan or viral infections, CC inflammation, allergies, Crohn's disease, diabetes, graft versus host CC disease, Parkinson's disease, multiple sclerosis, pain, psoriasis, CC anxiety, depression, schizophrenia, dementia, mental retardation, memory CC loss, epilepsy, asthma, tuberculosis, obesity, nausea, hypertension, CC any other disease, and immunodiagnosis. Abz42523 to Abz42869 encode CC exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G protein-coupled receptor; GPCR; antigenic peptide; gene therapy; G protein-coupled receptor modulator; antibody; immune-related disease; growth-related disease; cell regeneration-related disease; AIDS; cancer; immunological-related cell proliferative disease; autoimmune disease; Alzheimer's disease; atherosclerosis; infection; osteoarthritis; allergy; osteoporosis; cardiomyopathy; inflammation; Crohn's disease; diabetes; graft versus host disease; Parkinson's disease; multiple sclerosis; pain; psoriasis; anxiety; depression; schizophrenia; dementia; memory loss; mental retardation; epilepsy; asthma; tuberculosis; obesity; nausea; hypertension; hypotension; renal disorder; rheumatoid arthritis; trauma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 1; 523pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated antigenic peptides e.g., for G protein-coupled receptors (GPCR), useful for diagnosing and designing drugs for treating conditions in which GPCRs are involved, e.g. AIDS, Alzheimer's disease, cancer or
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                                          exemplification
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                                             invention
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Query Match Best Local Similarity Matches 317; Conser

99.8%; ilarity 99.7%; Conservative

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9; DB 6; Length 318; 2.2e-169;

Indels

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Sequence 318

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RESULT 20
ADB14370
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09-FEB-1998;
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14-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human prostate specific protein P835P
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N-PSDB; ADB14366, ADB14368, ADB14369.
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09-MAY-2001; 2001US-00852911.
29-JUN-2001; 2001US-00895814.
10-DEC-2001; 2001US-00012896.
09-MAY-2002; 2002US-00144678.
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US-10-001-469-288-3
US-10-001-469-288-3
US-10-001-469-288-3
US-10-147-368-3
US-10-10-43-527
US-09-86-055-81
US-09-968-055-81
US-09-968-055-81
US-10-10-144-678A-527
US-10-10-144-678A-527
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US-10-10-144-678A-527
US-10-10-10-144-633-65
US-10-10-10-144-633-65
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US-10-10-10-144-633-65
US-10-10-10-1469-288-12
US-10-968-033C-2
US-10-10-10-1-66-38
US-10-10-10-66-4
US-10-10-10-3-60-1-8
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2884, Ap
2884, Ap
2886, Ap
2886
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ALIGNMENTS

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Sequence 2880, Application US/10001469
; Sequence 2880, Application No. US20030091562A1
; GENERAL INFORMATION:
APPLICANT: PAROFONITS, AYA
APPLICANT: RAITANO, ARTHUR
APPLICANT: AFAR, DANIEL
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: FARIS, MARY
APPLICANT: FARIS, MARY
APPLICANT: OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 101P3A11 USEFUL IN TREATMENT AND DETECTION OF CANCE
FILIE OF INVENTION NUMBER: US/10/001,469
CURRENT APPLICATION NUMBER: 60/157,902
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR APPLICATION NUMBER: 60/291,118
PRIOR APPLICATION NUMBER: 60/291,118
PRIOR APPLICATION NUMBER: 09/680,728
PRIOR APPLICATION NUMBER: 09/680,728
PRIOR APPLICATION NUMBER: 09/680,728
PRIOR FILING DATE: 2000-10-05
SEQ ID NO 2880
LENGTH: 318
TYDER: DRT

TYPER: DRT

RESULT 2
US-10-147-368-19
; Sequence 19, Application US/10147368
; Publication No. US20030213004A1
, GENERAL INFORMATION:
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Best Local S
Matches 318
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Similarity 100.0%; Pred. No. 1.7e-151;
18; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                  IRQRILRLFHVATHASEP 318
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Sequence 28, Application US/10147368
Publication No. US20030213004A1
GENERAL INFORMATION:
APPLICANT: Agenays, Inc.
APPLICANT: Aya Jakobovits
APPLICANT: Mary Faris
APPLICANT: Mary Faris
APPLICANT: Robert Kendall Morrison
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran

Mary Faris Arthur B. Raitano Robert Kendall Morrison Douglas Saffran Wangmao Ge

APPLICANT

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APPLICANT: Ag Jakobovits
APPLICANT: Mary Faris
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Arthur B. Raitano
APPLICANT: Douglas Saffran
APPLICANT: Douglas Saffran
APPLICANT: Wangmao Ge
APPLICANT: Wangmao Ge
TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS
TITLE OF INVENTION: DETECTION OF CANCER
TITLE OF INVENTION: DETECTION OF CANCER
TITLE OF INVENTION: DETECTION OF CANCER
FILE REFERENCE: 51158-20024.21
CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: US 10/017,066
PRIOR APPLICATION NUMBER: US 10/017,066
PRIOR APPLICATION NUMBER: US 10/017,066
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR FILING DATE: 2001-05-5
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR APPLICATION NUMBER: US 09/680,728
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RESULT 3
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US-10-147-368-19
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LENGTH: 318
TYPE: PRT
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                          IRQRILRLFHVATHASEP 318
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PRIOR APPLICATION NUMBER: US 10/017,066
PRIOR APPLICATION NUMBER: US 10/001,469
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR PILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-05-15
PRIOR PILING DATE: 2001-05-15
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR PILING DATE: 1999-10-05
PRIOR 
                                                   Sequence 920, Application US/09759143
PREENT NO. US20022248A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Henderson, Robert A.
APPLICANT: Kalos, Michael D.
APPLICANT: Fanger, Gary R.
APPLICANT: Retter, Marc W.
APPLICANT: Retter, Marc W.
APPLICANT: Scolk, John A.
APPLICANT: Day, Craig H.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Vedvick, Thomas S.
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TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS
TITLE OF INVENTION: ENTITLED 10193A11 or PHOR-1 USEFUL IN TREATMENT
TITLE OF INVENTION: DETECTION OF CANCER
FILE REFERENCE: 51158-20024.21
CURRENT APPLICATION NUMBER: US/10/147,368
CURRENT FILING DATE: 2002-12-16
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Jiang, Yuqui
Henderson, Robert A.
Kalos, Michael D.
Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKE 300
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; APPLICANT: Wang, Aljun
; APPLICANT: Skeiky, Yasir A.W.
; APPLICANT: Skeiky, Yasir A.W.
; APPLICANT: Hepler, William
; TITLE OP INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OP INVENTION: DIAGNOSIS OF PROSTATE CANCER
; TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; TITLE OP INVENTION: DIAGNOSIS OF PROSTATE CANCER
; TITLE OP INVENTION NUMBER: US/09/759,143
; CURRENT FILING DATE: 2001-01-12
; NUMBER OF SEQ ID NOS: 934
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 920
; SEQ ID NO 920
; SEQ ID NO 920
; LENGTH: 318
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-759-143-920

99.8%; Score 1619; DB 3; Length 318;
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                                                       APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 920, Application US/09780669 Patent No. US20020051977A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 317;
APPLICANT: MCNeill, Patricia D.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C24
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Henderson, Robert /
Kalos, Michael D.
Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
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Hepler, William
                                                                                              Hural, John
                                                                                                                                                                                              Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
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Dillon, Davin C.
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99.7%; Pred. No. 3.4e-151;
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CURRENT APPLICATION NUMBER: US/09/780,669; CURRENT FILING DATE: 2001-02-09; NUMBER OF SEQ ID NOS: 943; SOFTWARE: FRatiseQ for Windows Version 3.0; SEQ ID NO 920; SEQ ID NO 920; LENGTH: 318; TYPE: PRT; ORGANISM: Homo sapiens
US-09-780-669-920
                                                                                                                                                                                                                                                                                                                         ; Sequence 920, Application US/09822827
; Patent No. US20020081680A1
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; FILE REFERENCE: 210121.534C1
; CURRENT APPLICATION NUMBER: US/09/822,827
; CURRENT FILING DATE: 2001-03-28
; NUMBER OF SEQ ID NOS: 982
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 920
; LENGTH: 318
; TYPE: PRT
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Best Local Similarity
Matches 317; Conserv
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)9-822-827-920
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Pred. No. 3.4e-151;
1; Mismatches 0;
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US-09-895-793-920
Sequence 920, Application US/09895793
Publication No. US20020192763A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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Patent No. US20020132273A1
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APPLICANT: ZOZULYA, SERGEY
TITLE OF INVENTION: RECEPTOR FINGERPRINTING, SENSORY PERCEPTION, AND
TITLE OF INVENTION: BIOSENSORS OF CHEMICAL SENSANTS
FILE REFERENCE: 078003-0277150
CURRENT APPLICATION NUMBER: US/9/886,055
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: 60/213,812
PRIOR APPLICATION NUMBER: 60/213,812
PRIOR FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 522
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TYPE: PRT
ORGANISM: Homo sapiens
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IRORILRLFHVATHASEP
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Sequence 920, Application US/0989581
Publication No. US20020193296A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Jiang, Yuqiu
APPLICANT: Kalos, Michael D.
APPLICANT: Retter, Marc W.
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REPERENCE: 210121.534C2
CURRENT APPLICATION NUMBER: US/09/895,793
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
TYPE: PRT
ORGANISM: Homo sapiens
US-09-895-793-920
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Skeiky, Yasir A.W.
Hepler, William T.
Henderson, Robert A.
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Carter, Darrick
Li, Samuel X.
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Stolk, John A.
Day, Craig H.
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Houghton, Raymond L.
Vinals de Bassols, Carlota
                                                                                                            Application US/09895814 o. US20020193296A1
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Kalos, Michael D.
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Pred. No. 3.4e-151;
1; Mismatches 0; Indels
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Sequence 2, Application US/09966459A

Publication No. US20030022237A1

GENERAL INFORMATION:
APPLICANT: FEDER, J. N.
APPLICANT: RAMANATHAN, C.S.
APPLICANT: HAWKEN, D.R.
APPLICANT: CACACE, A.
APPLICANT: CACACE, A.
APPLICANT: KORNACKER, M.G.
TITLE OF INVENTION: A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGP
TITLE OF INVENTION: EXPRESSED HIGHLY IN PROSTATE, COLON, AND LUN;
FILE REFERENCE: D0039NP
CURRENT APPLICATION NUMBER: 116/00/00/
                                                                                                                                                                                                                                                                                       RESULT 10
US-09-966-459A-2
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US-09-895-814-920
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C26
CURRENT APPLICATION NUMBER: US/09/895,814
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 990
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
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Best Local Similarity 99.7%;
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Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
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Vedvick, Thomas
Carter, Darrick
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Thomas
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Pred. No. 3.4e-151;
1; Mismatches 0;
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TITLE OF INVENTION. HUMAN OLFACTORY RECEPTORS AND GENES ENCODING SAME FILE REFERENCE: P 0278005
CURRENT APPLICATION NUMBER: US/09/804,291
CURRENT PILING DATE: 2001-03-13
PRIOR FILING DATE: 2000-03-13
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 2000-04-14
PRIOR FILING DATE: 2000-04-14
PRIOR FILING DATE: 2000-04-14
PRIOR FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/199,335
PRIOR APPLICATION NUMBER: 60/219,335
PRIOR APPLICATION NUMBER: 60/213,349
PRIOR FILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 60/213,849
PRIOR PILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/213,849
PRIOR APPLICATION NUMBER: 60/213,849
PRIOR APPLICATION NUMBER: 60/213,849
PRIOR APPLICATION NUMBER: 60/213,849
PRIOR APPLICATION NUMBER: 60/226,534
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; TYPE: PRT
; ORGANISM: Homo E
US-09-966-459A-2
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FRIOR APPLICATION NUMBER: 60/235,833
FRIOR FILING DATE: 2000-09-27
FRIOR APPLICATION NUMBER: 60/261,776
PRIOR APPLICATION NUMBER: 60/261,776
PRIOR APPLICATION NUMBER: 60/305,351
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: 60/313,202
PRIOR APPLICATION NUMBER: 60/313,202
PRIOR APPLICATION NUMBER: 60/313,202
PRIOR FILING DATE: 2001-08-17
NUMBER OF SEQ ID NOS: 60
SOPTWARE: PATENTIN Ver. 2.1
SEQ ID NO 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 91, Application US/09804291 Publication No. US20030088059A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 317; Conserv
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               FILING DATE: 2000-08-16
APPLICATION NUMBER: 60/230,732
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2000-09-07
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Pred. No. 3.4e-151;
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RESULT 12
US-10-012-896-920
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US-09-804-291-91
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Best Local Similarity
Matches 317; Conserv
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SOFTWARE: Patentin Ver. 2
SEQ ID NO 91
LENGTH: 318
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Publication No.
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Vināls de Bassols, Carlota
APPLICANT: Foy, Teresa
APPLICANT: Fanger, Garry R.
APPLICANT: Wantanabe, Yoshihiro
APPLICANT: Weagher, Madeleine Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
TILE REFERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
                                                                                                                                                                                      APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2001-02-07
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Stolk, John A.
Day, Craig H.
Vedvick, Thomas &
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Vedvick, Thomas
Carter, Darrick
                                                                                                                                                                  McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
                                                                                                                                                                                                                                                           Skeiky, Yasir A.W.
Hepler, William T.
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Kalos, Michael D.
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                                                                                                                                                                                                                                           Henderson, Robert
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5. US20020183251A1
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Pred. No. 3.4e
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CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/257,144
PRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
SEQ ID NO 434
LENGTH: 318
TYPE: PRT
ORGANISM: Homo sapiens
US-10-225-567A-434
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US-10-225-567A-434
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Best Local (
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Publication No. US20030113798A1
GENERAL INFORMATION:
APPLICANT: LifeSpan Biosciences
APPLICANT: Brown, Joseph P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 99.7
Matches 317; Conservative
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                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 1920-4-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo gapiens
-10-012-896-920
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Pred. No. 3.4e-151;
1; Mismatches 0;
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APPLICANT: AGAI, KIYOSHI
APPLICANT: AGAI, KIYOSHI
APPLICANT: AGAI, KIYOSHI
APPLICANT: AKURAMAI, HIROYUKI
ITILE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT TLING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR PILLNG DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 868
RESULT 15
US-10-144-678A-920
US-10-144-678A-920
; Sequence 920, Application US/10144678A
; Publication No. US20030157089A1
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
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Best Local Similarity 99.7
Matches 317; Conservative
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TYPE: PRT
ORGANISM: Homo sapiens
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APPLICANT:

Henderson, Robe Kalos, Michael

Yugiu

Susan L. iu Robert A.

Dillon, Davin C. Mitcham, Jennifer

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APPLICANT: Watanabe, Yoshihiro

APPLICANT: Deng, Ta

APPLICANT: Deng, Ta

APPLICANT: Deng, Ta

APPLICANT: Deng, Ta

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 210121.427C28

CURRENT APPLICATION NUMBER: US/10/144,678A

CURRENT FILING DATE: 2002-08-12

NUMBER OF SEQ ID NOS: 1033

SOFTWARE: FASTSEQ for Windows Version 3.0

SEQ ID NO 920

LENGTH: 318

TYPE: PRT

ORGANISM: Homo sapiens

US-10-144-678A-920
                                                                                                      RESULT 16
US-10-294-025-920
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Sequence 920, Application US/10294025
Publication No. US20030185830A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Kalos, Michael D.
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                                                                                                                                                                      IRQRILRLFHVATHASEP
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Hepler, William T.
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Carter, Darrick
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APPLICANT: Wernet, Kumud
APPLICANT: Vernet, Corine
APPLICANT: Casman, Stacie J
APPLICANT: Casman, Stacie J
APPLICANT: Casman, Stacie J
APPLICANT: Wolenc, Adam R
APPLICANT: Boderna, Steven K
APPLICANT: Hishnu, Vishun S
APPLICANT: Hishnu, Vishun S
APPLICANT: Spytek, Kimberly A
APPLICANT: Syytek, Kimberly A
APPLICANT: Gusev, Valdimir
TITLE OF INVENTION: NO. US20030195335A1el Proteins and Nucleic Acid
FILLE REFERENCE: 15966-748
CURRENT APPLICATION NUMBER: US/10/044,643
CURRENT FILING DATE: 2002-01-11
PRIOR APPLICATION NUMBER: 60/193,664
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: 60/194,614
PRIOR APPLICATION NUMBER: 60/194,614
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: 60/195,063
PRIOR APPLICATION NUMBER: 60/195,063
PRIOR APPLICATION NUMBER: 60/195,066
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: 60/195,067
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FILE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER |
FILE REFERENCE: 210121.427C29 |
CURRENT APPLICATION NUMBER: US/10/294,025 |
CURRENT FILING DATE: 2002-11-12 |
NUMBER OF SEQ ID NOS: 1038 |
SOFTMARE: FastSEQ for Windows Version 3.0 |
SEQ ID NO 920 |
SEQ 
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US-10-044-643-26
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Pred. No. 3.4e-151;
1; Mismatches 0;
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Same

APPLICATION NUMBER: 60/195,068 FILING DATE: 2000-04-06

APPLICATION NUMBER: 60/195,069

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RESULT 18
US-10-147-368-32
JSequence 32, Application US/10147368
Publication No. US20030213004A1
GENERAL INFORMATION:
APPLICANT: Agensys, Inc.
APPLICANT: Aya Jakobovits
APPLICANT: Arthur B. Raitano
APPLICANT: Robert Kendall Morrison
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PRIOR FILING DATE: 2001-01-30
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 26
LENGTH: 318
TYPE: PRT
TYPE: PRT
TORGANISM: Homo sapiens
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FILING DATE: 2000-07-27
APPLICATION NUMBER: 60/221,325
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APPLICATION NUMBER: 60/224,588
FILING DATE: 2000-08-11
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APPLICATION NUMBER: 60/219,855
FILING DATE: 2000-07-21
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APPLICATION NUMBER: 60/195,070
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Pred. No. 3.4e-151;
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APPLICANT: HAGA, TATSUYA
FITTLE OF INVENTION: NOVEL G-PROTEIN COUPLED REF
FILE REFERENCE: 31671-186347
CURRENT APPLICATION UNMER: US/10/343,650A
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION UNMER: JP 2000/237818
PRIOR PILING DATE: 2000-08-04
PRIOR APPLICATION UNMER: JP 2001/34434
PRIOR FILING DATE: 2001-02-13
NUMBER OF SEQ ID NOS: 694
SOFTWARE: PATENTIN Ver. 2.1
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APPLICANT: Wangmao Ge
APPLICANT: Pia M. Challita-Eid
TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS
TITLE OF INVENTION: ENTITLED 10193A11 or PHOR-1 USEFUL IN 7
TITLE OF INVENTION: ENTITLED 10193A11 or PHOR-1 USEFUL IN 7
TITLE OF INVENTION: DETECTION OF CANCER
FILE REFERENCE: 51158-20024.21
CURRENT APPLICATION NUMBER: US/10/147,368
CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: US 10/017,066
PRIOR FILING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US 10/001,469
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR APPLICATION NUMBER: US 60/291,118
PRIOR APPLICATION NUMBER: US 60/291,728
PRIOR PILING DATE: 2001-05-15
PRIOR APPLICATION NUMBER: US 60/157,902
PRIOR PILING DATE: 2000-10-05
PRIOR PILING DATE: 1999-10-05
PRIOR PILING DATE: 1999-10-05
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US-10-343-650A-158
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Best Local Similarity 99.7%;
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TYPE: PRT
ORGANISM: Homo 9
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Pred. No. 3.4e-151;
1; Mismatches 0;
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Sequence 10, Application US/10692605
; Publication No. US20040091928A1
; GENERAL INFORMATION:
; APPLICANT: Vogeli, Gabriel
; APPLICANT: Wood, Linda S.
; APPLICANTION UMMBER: US/10/692,605
; CURRENT FILING DATE: 2000-10-27
; PRIOR APPLICATION NUMBER: US 09/481,794
; PRIOR APPLICATION NUMBER: US 09/481,794
; PRIOR APPLICATION NUMBER: US 09/454,399
; PRIOR APPLICATION NUMBER: US 09/454,399
; PRIOR APPLICATION NUMBER: US 09/459,517
; PRIOR APPLICATION NUMBER: US 09/429,517
; PRIOR APPLICATION NUMBER: US 09/429,555
; PRIOR FILING DATE: 1999-10-28
; PRIOR APPLICATION NUMBER: US 09/429,555
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US-10-692-605-10
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-343-650A-158
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                                       Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 75
SOPTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 318
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Best Local Similarity 99.7%;
Matches 317; Conservative
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PRIOR APPLICATION NUMBER: US 09/428,020
PRIOR FILING DATE: 1999-10-27
PRIOR APPLICATION NUMBER: US 09/427,859
PRIOR FILING DATE: 1999-10-27
ORGANISM: Homo sapiens
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APPLICATION NUMBER: US 09/429,676
FILING DATE: 1999-10-28
APPLICATION NUMBER: US 09/429,695
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APPLICATION NUMBER: US 09/428,114
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Pred. No. 3.4e-151;
1; Mismatches 0;
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Search completed: March
Job time : 166 secs
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Best Local S
Matches 317
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61 PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQWFAIHSLSGMESTVLLA
61 PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQWFAIHSLSGMESTVLLA
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317; Conserv
                                                                        IRQRILRLFHVATHASEP 318
                                                                                                   GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKE 300
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99.7%;
             2006, 08:43:56
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Pred. No. 3.4e-151;
1; Mismatches 0;
                                                    318
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2322001154
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3: /cgn2-6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

4: /cgn2-6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*

5: /cgn2-6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*

6: /cgn2-6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*

7: /cgn2-6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*

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US-10-055-877-335

US-11-190-188-10

US-11-190-188-9

US-11-190-188-9

US-11-124-367A-496

US-11-124-367A-514

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US-10-055-877-337

US-10-055-877-222

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US-10-055-877-234
US-11-191-188-16
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US-11-191-188-18
US-11-191-188-19
US-10-11-191-188-19
US-11-191-188-19
US-11-191-1
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-234-786-527
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SOFTWARE: FAST:
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FONGTH: 320
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Sequence 527, Application US/11234786
Publication No. US20060024301A1
GENERAL INFORMATION:
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TITLE OF INVENTION: PROSTATE-SPECIFIC POLYPEPTIDES AND FUSION
TITLE OF INVENTION: POLYPEPTIDES THEREOF
FILE REFERENCE: 210121.427C31
CURRENT APPLICATION NUMBER: US/11/234,786
CURRENT FILING DATE: 2005-09-23
PRIOR APPLICATION NUMBER: US 09/568,857
PRIOR FILING DATE: 2000-05-09
PRIOR FILING DATE: 2000-05-07
PRIOR APPLICATION NUMBER: US 09/536,857
PRIOR APPLICATION NUMBER: US 09/536,857
PRIOR FILING DATE: 2000-05-27
PRIOR FILING DATE: 2000-01-14
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APPLICATION NUMBER: US 09/030,607
FILING DATE: 1998-02-25
ning Prior Application data removed -
R OF SEQ ID NOS: 701
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FILING DATE: 1999-04-09
APPLICATION NUMBER: US 09/232,149
FILING DATE: 1999-01-15
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FILING DATE: 1999-11-12
APPLICATION NUMBER: US 09/352,616
FILING DATE: 1999-07-13
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APPLICATION NUMBER: US 09/115,453
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Reed, Steven G.
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Vedvick, Thomas S.
Carter, Darrick
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Mitcham, Jennifer L.
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55; Mismatches 67
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   123
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                                            APPLICANT: Casman, Stacie
APPLICANT: Boldog, Ference
TITLE OF INVENTION: NOVAL Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2001-01-2
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-2
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,119
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,399
PRIOR FILING DATE: 2001-01-25
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PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,391
PRIOR APPLICATION NUMBER: 60/263,351
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PRIOR APPLICATION NUMBER: 60/272,870
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APPLICATION NUMBER: 60/272,870 FILING DATE: 2001-03-02 APPLICATION NUMBER: 60/275,990
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Spytek, Kimberly
Ratelli, Luca
Kekuda, Ramesh
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Zerhusen, Bryan
Andrew, David
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Miller, Charles
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Baumgartner, Jason
Shimkets, Richard
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Taupier Jr., Raymond
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Burgess, Cahterine
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o. US20050288241A1
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; PRIOR FILING DATE: 2001-03-14
; PRIOR APPLICATION NUMBER: 60/275,927
; PRIOR FILING DATE: 2001-03-14
; Remaining Prior Application data removed - Sei NUMBER OF SEQ ID NOS: 512
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 336
; ERNOTH: 329
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-336
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Sequence 339, Application US/10055877
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
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Wolenc, Adam
Baumgartner, Jason
Shimkete, Richard
Gusev, Vladimir
Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
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Zerhusen, Bryan
Andrew, David
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Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
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Patturajan, Meera
Burgess, Cahterine
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Miller, Charles
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Kekuda, Ramesh
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Pred. No. 2.5e-74;
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PRIOR PILLING DATE: 2001-01-23
PRIOR PILLING DATE: 2001-01-24
PRIOR PILLING DATE: 2001-01-24
PRIOR PILLING DATE: 2001-01-24
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-25
PRIOR PILLING DATE: 2001-01-35
PRIOR PILLING DATE: 2001-01-30
PRIOR PILLING DATE: 2001-01-30
PRIOR PILLING DATE: 2001-03-14
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 339
LENGTH: 312
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Best Local :
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CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
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APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
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305
                                                                                     245
                                                                                                                    247 VCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKBIRQRIL 306
                                                                                                                                                                                                           188 MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLG-LTREAQAKAFGTCVSH
                                                                                                                                                                                                                                                                       125 AICDPLRYATVLTTEVIAAMGLGAAARSFITLFPLPFLIKRLPICRSNVLSHSYCLHPDM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   163;
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                                        RLFH 310
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RMFH 308
                                                                                     ILAVLAFYVPMIGVSTVHRFGKHVPCYIHVLMSNVYLFVPPVLNPLIYSAKTKEIRRAIF
                                                                                                                                                                               MRLACADISINSIYGLFVLVSTFGMDLFFIFLSYVLILRSVMATASREERLKALNTCVSH
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53.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 868.5; DB 6;
Pred. No. 3.7e-72;
1; Mismatches 89;
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; Sequence 8, Application US/11190188; Publication No. US20060035331A1; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.; APPLICANT: LAL, Preeti; APPLICANT: TANG, Y. Tom
; APPLICANT: PATTERSON, Chandra

RESULT 4 US-11-190-188-8

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RESULT 5
US-10-511-538-70
VS-10-511-538-70
Sequence 70, Application US/10511538
Publication No. US20060026700A1
GENERAL INFORMATION:
APPLICANT: OriGene Technologies, Inc
TITLE OF INVENTION: TISSUE SPECIFIC GENES AN
FILE REFERENCE: 16U 200 PCT
CURRENT APPLICATION NUMBER: US/10/511,538
CURRENT FILING DATE: 2004-10-18
PRIOR APPLICATION NUMBER: US 60/372,669
PRIOR FILING DATE: 2002-04-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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APPLICANT:
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APPLICANT: BOROWSKY, Mark L.
TITLE OP INVENTION: G-PROTEIN COUPLED RECEPTORS
FILE REFERENCE: PI-0044 PCT
CURRENT APPLICATION NUMBER: US/11/190,188
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PRIOR APPLICATION NUMBER: US/10/220,382
PRIOR FILING DATE: 2001-03-01
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NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No: 7472361CD1
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TYPE: PRT
ORGANISM: Homo sapiens
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Local Similarity 49.8%; Pred. No. 2.1e-68;
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AU-YOUNG, Janice
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                                                                                                  AND
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; TYPE: PRT
; ORGANISM: Homo s
US-10-511-538-70
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Best Local Similarity 50.7:
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 335, A Publication No.
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OR APPLICATION NUMBER: US 60/424,336
OR FILING DATE: 2002-11-07
OR APPLICATION NUMBER: US 60/374,823
OR FILING DATE: 2002-04-24
OR APPLICATION NUMBER: US 60/376,558
OR FILING DATE: 2002-05-01
OR APPLICATION NUMBER: US 60/381,366
OR APPLICATION NUMBER: US 60/403,648
OR APPLICATION NUMBER: US 60/403,648
OR FILING DATE: 2002-08-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 60/411,882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            133 LRHATVLTLERVTKIGVAAVVRGAALMAPLEVFIKQLEFCRSNILSHSYCLHQDVMKLAC 192
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Gerlach, Valerie
Spytek, Kimberly
Ratelli, Luca
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                           Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
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Miller, Charles
                                                                                                                   Mezes, Peter
Patturajan, Meera
Burgess, Cahterine
           Vernet,
                                                                              Wolenc, Adam
                                                                                                Eisen, Andrew
                                                                                                                                                                  Andrew, David
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Zerhusen, Bryan
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b. US20050288241A1
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Indels Length

Gaps

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314; 1;

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RESULT 7
US-11-190-188-10
; Sequence 10, Application US/11190188
; Publication No. US20060035331A1
; GENERAL INFORMATION:
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US-10-055-877-335
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NUMBER OF SEQ ID NOS: 512
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 335
LENGTH: 342
TYPE: PRT
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
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TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids
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APPLICATION NUMBER: 60/264,478
FILING DATE: 2001-01-26
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FILING DATE: 2001-01-24
APPLICATION NUMBER: 60/264,117
FILING DATE: 2001-01-25
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APPLICATION NUMBER: 60/275,990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
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SHISAVSIFYLPLISLSLVHRYGHSAPPFVHIIMANVFLLIPPVLNPIIYSVKIKQIQKA 316
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                                                                                                                  IIKV 320
                                                                                                                                              ILRL 308
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46.7%; Pred. No. 8.4e-66;
ative 72; Mismatches 89;
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APPLICANT:

Ballinger, Robert

Anderson, David

Padigaru, Muralidhara Miller, Charles Tchernev, Velizar

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Sequence 338, Application US/100
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidha
APPLICANT: Miller, Charles
APPLICANT: Tchernev, Velizar
APPLICANT: Zhong, Mei
                                                                                                                                                             RESULT 8
US-10-055-877-338
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CURRENT FILING DATE: 2005-07-27
PRIOR APPLICATION NUMBER: US/10/220,382
PRIOR FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: 60/186,854; 60/188,384; 60/190,453; 60/190,730
PRIOR PILING DATE: 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PERL Program
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 47.5%; Score 771; DB 7; Length Best Local Similarity 47.3%; Pred. No. 3.1e-63; Matches 140; Conservative 71; Mismatches 83; Indel
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                134 YATILTSLVLVGISMCIVIRPVLLTLPMVYLIYRLPFCQAHIIAHSYCEHMGIAKLSCGN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           74 ALSATSVPRMLGIFWFDAHEINYGACVAQMFLIHAFTGMEAEVLLAMAFDRYVAICAPLH 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                75 LISTSSMPKMLAIFWENSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYVAICHPLR 134
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                                                                                                                                                                                                                                             YIPSVFSFLTHRFGHQIPGYIHILVANLYLIIPPSLNPIIYGVRTKQIRERVLYVF 308
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TRIBOULEY, Catherine
LU, Dyung Aina M.
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TANG, Y. To
                                                                                                                       Application US/10055877
o. US20050288241A1
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SOPTWARE: Pate
SEQ ID NO 338
FONCTH: 321
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Best Local Similarity 45.2%;
Matches 140; Conservative 73
                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens -10-055-877-338
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CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR FILING DATE: 2001-01-19
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TYPE: PRT
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TITLE OF INVENTION: Novel Polypeptides and Nucleic
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APPLICATION NUMBER: 60/264,117
FILING DATE: 2001-01-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/263,351 FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/264,139 FILING DATE: 2001-01-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2001-03-1
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APPLICATION NUMBER: 60/275,990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
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                    181
                                                                         121 MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS 180
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YCLHQDVMKLACDDIRVKYVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKA 239
                                                                                                                                                                                                  MITSVSPSTSTNSSPLLTGFSGMEQQYPWFSIPFSSIYAMVLLGNCMVLHVIWTEPSLHQ 66
                                                                                                                              PMFYFLSMLALTDLCMGLSTVYTVLGILWRIIREISLDSCIAQSYFIHGLSFMESSVLLT 126
                                                                                                                                                                  PMYIPLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
                                                                                                                                                                                                                                      MWVDPNGNESSATYFILIGLPGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHE 60
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Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
                                                        MAFDRYIAICNPLRYSSILTNSRIIKIGLTIIGRSFFFITPPIICLKFFNYCHFHILSHS 186
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Taupier Jr., Raymond
Pena, Carol
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Burgess, Cahterine
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Mezes, Peter
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Guo, Xiaojia
Tarhusen, Bryan
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Spytek, Kimberly
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Pred. No. 4.4e-62;
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; ORGANISM: Homo sapiens
; FRATURE;
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No: 7472363CD1
US-11-190-188-9
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US-11-190-188-9
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45.7%; Score /**, ---
Best Local Similarity 46.7%; Pred. No. 1.6e-6
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CURRENT APPLICATION NUMBER: US/11/190,188
CURRENT FILING DATE: 2005-07-27
PRIOR APPLICATION NUMBER: US/10/220,382
PRIOR FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: 60/186,854; 60/188,384; 60/190,453; 60/190,730
PRIOR FILING DATE: 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PERL Program
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TYPE: PRT
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INVENTION: G-PROTEIN COUPLED RECEPTORS
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SMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKBIRQRILRLF 309
                                                           GLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSHVCAVFIFYVPFIGL
                                                                                                                                                           GLEDFHMWISGPFCSVYLVALLGNATILLVIKVEQTLREPMFYFLAILSTIDLALSATSV
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                                 GLFV-VSFFVLNLVLIGISYVYILRAVFRLPSHDAQLKALSTCGAHVGVICVFYIPSVFS
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TRIBOULEY, Cather
LU, Dyung Aina M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Terence P.
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Pred. No. 1.6e-60;
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                                                                                                                                                                                                                                                                                                                       Length 356;
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DB 7; 93;

Indels Length 312;

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; ORGANISM: Homo sapiens
US-11-124-367A-496
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US-11-124-367A-496
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Publication No. US20060024700A1
GENERAL INFORMATION:
APPLICANT: Michele Cargill
APPLICANT: Hongjin Huang
TITLE OP INVENTION: Generatic Polymorphisms Associated with
TITLE OP INVENTION: Fibrosis Methods of Detection and Uses Thereof
FILE REFERENCE: CL001519.ORD
                                                                                                                                                            Sequence 514, Application US/11124367A Publication No. US20060024700A1 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 34460
SOPTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 496
LENGTH: 312
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Best Local (
APPLICANT: Michele Cargill
APPLICANT: Hongjin Huang
TITLE OF INVENTION: Genetic Polymorphisms Associated with
TITLE OF INVENTION: Fibrosis Methods of Detection and Uses
FILE REFERENCE: CL001519.ORD
CURRENT APPLICATION NUMBER: US()11/124,367A
CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US 60/568,846
PRIOR FILING DATE: 2004-05-07
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CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US 60/568,846
PRIOR FILING DATE: 2004-05-07
PRIOR APPLICATION NUMBER: US 60/582,609
PRIOR PILING DATE: 2004-06-25
PRIOR APPLICATION NUMBER: US 60/599,554
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   62 AMLAATDLGVTLTIMPTVLGVLWLDHREIGHGACFSQAYFIHTLSVMBSGVLLAMAYDCF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67 CMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAMAFDRY 126
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Pred. No. 3.6e-60;
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PRIOR FILING DATE: 2004-06-25
PRIOR PILING DATE: 2004-08-09
PRIOR FILING DATE: 2004-08-09
NUMBER OF SEQ ID NOS: 34460
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 514
PRIOR THE STATE OF SEC ID NOS: 34460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 123, Appropriation No.
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Best Local Similarity 45.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       186 DVMKLACDDIRVNVVYGLIVIISAIGLDSLLISPSYLLILKTVLGL-TREAQAKAFGTCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         128 AICHPLRHATVLTLPRVTKIGVAAVVRGAALMAP--LPVFIKQLPFCRSNILSHSYCLHQ 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             63 MLAATDLGLALTIMPTVLGVLWLDHREIGSAACFSQAYFIHSLSFLESGILLAMAYDRFI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68 MLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYV 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 SSSSSHPFLLTGFPGLEEAHHWISVFFLFMYISILFGNGTLLLLIKEDHNLHEFMYFFLA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKEIRQR 304
                                                                                                                                                                                                                                                                                                                        Anderson, David
Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
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Miller, Charles
                                                                     Shimkets, Richard
Gusev, Vladimir
Vernet, Corine
                                                                                                                                                                                                                 Andrew, David
Mezes, Peter
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Zerhusen, Bryan
 Shenoy, Suresh
Li, Li
                                                                                                                          Wolenc, Adam
Baumgartner, Jason
                                                                                                                                                              Burgess, Cahte
Bisen, Andrew
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                                   Pena,
                                                    Taupier Jr., Raymond
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Burgess, Cahterine
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Kekuda, Ramesh
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b. US20050288241A1
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Pred. No. 4.4e-60;
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US-10-055-877-337; Sequence 337, Application US/10055877; Publication No. US20050288241A1
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                                                          GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tcherney, Velizar
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SEQ ID NO 123
LENGTH: 303
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                                   APPLICANT:
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ORGANISM: Homo
-10-055-877-123
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PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
   APPLICANT:
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CURRENT FILING DATE: 2002-01-22
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APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/264,478 FILING DATE: 2001-01-26 APPLICATION NUMBER: 60/263,351
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FILING DATE: 2001-01-24
APPLICATION NUMBER: 60/264,117
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APPLICATION NUMBER: 60/264,139
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Zhong, Mei
Anderson, David
Ballinger, Robert
Gerlach, Valerie
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                                                                                                                                                                                                                                                                                                                                                                           IRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSHVCAVFIF 253
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Pred. No. 3.5e-59;
8; Mismatches 98
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; ORGANISM: Homo sapiens US-10-055-877-337
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CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR PILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR PILING DATE: 2001-01-24
PRIOR PILING DATE: 2001-01-24
PRIOR PILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR PILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR PILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR PILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR PILING DATE: 2001-01-30
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 337
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Best Local (
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APPLICANT: Shenoy, Suresh
APPLICANT: Li, Li
APPLICANT: Casman, Stacie
APPLICANT: Boldog, Ference
APPLICANT: Boldog, Ference
APPLICANT: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
FILE REFERENCE:
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TYPE: PRT
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FILING DATE: 2001-03-02
APPLICATION NUMBER: 60/275,990
FILING DATE: 2001-03-14
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                                                                                 186 DVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCV 244
                                                                                                                                                                      125
                                                                                                                                                                                                                                                   126 YVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQ 185
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EVIKYTYSKPWISSFWGLFLQLYLNGTDVLFILFSYVLILRTVLGIVARKKQQKALSTCV
                                                                                                                                                                      FVAICNPLNYATILTDRMVLVIGLVICIRPAVFLLPLLVAINTVSFHGGHELSHPFCYHP
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Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
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Taupier Jr., Raymond
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Mezes, Peter
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Burgess, Cahterine
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Zerhusen, Bryan
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Pred. No. 2.1e-58;
8; Mismatches 104;
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RESULT 15
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; ORGANISM: Homo sapiens
US-10-511-538-225
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US-10-511-538-225
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Best Local Similarity 41.3%; Pred. No. 5.1e-52;
Matches 126; Conservative 59; Mismatches 98;
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SEQ ID NO 225
LENGTH: 291
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NUMBER OF SEQ ID NOS: 344
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PRIOR FILING DATE: 2002-11-07
PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: US 60/374,823
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CURRENT APPLICATION NUMBER: US/10/511,538
CURRENT FILING DATE: 2004-10-18
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TITLE OF INVENTION: TISSUE SPECIFIC GENES AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/372,669
PRIOR FILING DATE: 2002-04-16
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                                                                                                                                                                                                                 245 SHVCAVEIFYVPFIGLSMYHRFSKRRDSPLFVILANIYLLVPFVLNPIVYGVKTKEIRQR 304
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                                                                                            ILLLF 286
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PRIOR FILING DATE: 2002-04-16
PRIOR PELICATION NUMBER: US 60/411,882
PRIOR FILING DATE: 2002-09-20
PRIOR APPLICATION NUMBER: US 60/424,336
PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR FILING DATE: 2002-04-24
                     APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tcherney, Velizar
APPLICANT: Zhong, Mei
APPLICANT: Anderson, David
APPLICANT: Ballinger, Robert
APPLICANT: Gerlach, Valerie
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Matches 114;
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                                                                                                                                                                                                       GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US 60/376,558

PRIOR FILING DATE: 2002-05-01

PRIOR APPLICATION NUMBER: US 60/381,366

PRIOR FILING DATE: 2002-05-20

PRIOR FILING DATE: 2002-05-20
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TITLE OF INVENTION: TISSUE SPECIFIC GENES
FILE REFERENCE: 16U 200 PCT
CURRENT APPLICATION NUMBER: US/10/511,538
CURRENT FILING DATE: 2004-10-18
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SOFTWARE: PatentIn version 3.1
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ORGANISM: Homo sapiens
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Anderson, David
Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
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TITLE OF INVENTION: Novel Polypeptides and Nucleic
FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR PILING DATE: 2001-01-23
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OR FILING DATE: 2001-01-24
OR APPLICATION NUMBER: 60/264,117
OR FILING DATE: 2001-01-25
OR APPLICATION NUMBER: 60/264,139
OR FILING DATE: 2001-01-25
                                                                                                                                                                                                                                                                                                                 / Match 29.3%; Score 475; DB 6; Length 318; Local Similarity 35.7%; Pred. No. 3.8e-36; ndels 114; Conservative 62; Mismatches 121; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/275,927
FILING DATE: 2001-03-14
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FILING DATE: 2001-03-02
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FILING DATE: 2001-01-30
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                                                                                                    RYVAICHPIRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPPCRSNILSHSYCLH 184
                                                                                                                                                                                                                                  NESSATYFILIGL---PGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYI 64
LAVLKLACADISINVISMGVANVIFLGVPVLFIFVSYIFILSTILRIPSAEGRKKAFSTC
                              QDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTC 243
                                                                        RYVAICNPLRYPVVMNKAAYVPMAVSSWVAGGANSLVQISLAVQLPPCGDNVINHFICEI 181
                                                                                                                                                        FLGNLSFLDICYTTSSVPLVLDGFLTPRKTISFSGCAVQMFLSFAMGATECVLLGMMAFD
                                                                                                                                                                                             FLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAMAFD 124
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Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
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Patturajan, Meera
Burgess, Cahterine
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Zerhusen, Bryan
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PILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/263,892
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-19
PRIOR PILING DATE: 2001-01-23
PRIOR PILING DATE: 2001-01-23
PRIOR PILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR APPLICATION NUMBER: 60/272,870
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/272,870
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR APPLICATION NUMBER: 60/275,990
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US-10-055-877-326
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Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 512
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 326
LENGTH: 318
TYPE: PRT
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PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
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Taupier Jr., Raymond
Pena, Carol
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Shimkets, Richard
Gusev, Vladimir
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Gerlach, Valerie
Spytek, Kimberly
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Miller, Charles
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Burgess, Cahterine
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Mezes, Peter
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Zerhusen, Bryan
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Kekuda, Ramesh
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b. US20050288241A1
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US-10-055-877-326
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Matches 114; Conservative
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APPLICANT:
APPLICANT:
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
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APPLICANT:
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                                                                           APPLICANT: Casman, Stacie
APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic
FILE REFERENCE: 21402-251
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Gerlach, Valerie
Spytek, Kimberly
Ratelli, Luca
Kekuda, Ramesh
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Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
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Taupier Jr., Raymond
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Burgess, Cahterine
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Mezes, Peter
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Miller, Charles
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                                                                                               Acids Encoded Thereby
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Sequence 12, Application US/11190188
Dilication No. US20060035331A1
GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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SHIH, Leo L.
TRIBOULEY, Catheri
LU, Dyung Aina M.
                                                                                                                                          TANG, Y. TO
PATTERSON,
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                KHAN, Farrah A.
POLICKY, Jennifer L.
AU-YOUNG, Janice
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PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR PELICATION NUMBER: 60/272,870
PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
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APPLICATION NUMBER: 60/264,139
FILING DATE: 2001-01-25
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               PRIOR APPLICATION NUMBER: US 60/411,882
PRIOR FILING DATE: 2002-09-20
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PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR APPLICATION NUMBER: US 60/376,558
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                                                                                                                                                                                                                                                                           Sequence 21, Application US/10511538
Publication No. US.20060026700A1
GENERAL INFORMATION:
APPLICANT: OriGene Technologies, Inc
TITLE OF INVENTION: TISSUE SPECIFIC GENES
FILE REFERENCE: 16U 200 PCT
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PRIOR PILING DATE: 2000-03-03; 2000-03-10; 2000-03-17; 2000-03-20
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PERL Program
SEQ ID NO 12
LENGTH: 319
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APPLICANT:
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ORGANISM: Homo sapiens
PEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyt
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CURRENT FILING DATE: 2004-10-18
PRIOR APPLICATION NUMBER: US 60/372,669
PRIOR FILING DATE: 2002-04-16
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PRIOR APPLICATION NUMBER: US/10/220,382
PRIOR FILING DATE: 2001-03-01
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CURRENT APPLICATION NUMBER: US/11/190,188
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APPLICANT: BOROWSKY, Mark L.
TITLE OF INVENTION: G-PROTEIN COUPLED RECEPTORS
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Pred. No. 3.8e-35;
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PRIOR APPLICATION NUMBER: US 60/403,648
PRIOR FILING DATE: 2002-08-16
NUMBER OF SEQ ID NOS: 344
SOFTWARE: Patentin version 3.1
SEQ ID NO 21
LENGTH: 319
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Best Local (
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                                                                                                                                                PVLSLACTOT-----FLVVIVDAIHAABIVASFLVIALSYIRIIIVILGMHSAEGHHKA
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DMKEAIGRLFHYQKRA
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Search completed: March Job time : 23 secs 9, 2006, 08:44:24

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odorant receptor 8
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                                     sulfakinin recepto gastric CCKA rece probable olfactory melanocortin receptor melanocortin receptor ladenosine receptor A(1) adenosine receptor A(1) adenosine receptor probable olfactory probable olfactory melanocortin receptor probable olfactory angiotensin I receptor ladenosine receptor brobable olfactory angiotensin I receptor probable olfactory angiotensin I receptor probable olfactory probable olfactory probable olfactory probable olfactory probable olfactory probable olfactory probable olfactory
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C;Species: Homo sapiens (man)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 26-Aug-1999
C;Accession: A60547; A60482
R;Feingold, E.A.; Forget, B.G.
Blood 74, 2178-2186, 1989
A;Title: The breakpoint of a large deletion causing hereditary persistence of fetal hemo A;Feference number: A60547; MUID:90028751; PMID:2478223
A;Accession: A60547
A;Accession: A60547
A;Residues: 1-292 <FEI>A;Rose-references: UNIFARC:UPF0000178388
C;Superfamily: olfactory receptor OR14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              odorant receptor 8 - channel catfish
(;Species: Ictalurus punctatus (channel catfish)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change
C;Accession: E45774
R;Ngai, J:; Dowling, M.M.; Buck, L.; Axel, R.; Chess, A.
Cell 72, 657-666, 1993
A;Title: The family of genes encoding odorant receptors in the caption of the caption 
                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q9PSJ5; UNIPARC:UPI0000042025
A;Experimental source: olfactory epithelium
A;Note: sequence extracted from NCBI backbone (NCBIP:1277.
C;Superfamily: olfactory receptor OR14
C;Keywords: olfaction; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Status: preliminary; not compared with conceptual translation A; molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein (HPFH breakpoint 3' region) - human N;Alternate names: olfactory receptor homolog
                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Residues: 1-313 <NGA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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Local Similarity 40.3%; Pred. No. 1.8e-40;
nes 114; Conservative 49; Mismatches 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 248
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                                                                                                                                                                                                Similarity
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                                     MLAPVQNISFTT-FTLTGFHDLGEWGPILSIPYLLMFLLSSTSNLTLIYLIISQRALHSP 59
                                                                                                                                                              29.2%; Score 473.5; DB 2; 34.1%; Pred. No. 1.2e-33; tive 66; Mismatches 132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ALIGNMENTS
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$29710
$29710
olfactory receptor OR18 - rat
C, Species: Rattus norvegicus (Norway rat)
C, Date: 25-Reb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Sep-1998
C; Accession: $29710
RESULT 4
A46247
Olfactory receptor OR3 - mouse
Olfactory receptor OR3 - mouse
C;Species: Mus musculus (house mouse)
C;Species: Mes musculus (house mouse)
C;Date: 22-Gep-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Date: 22-Gep-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: A46247
C;Accession: A46247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Raming, K.; Krieger, J.; Strotmann, J.; Boekhoff, I.; Kubick, Nature 361, 353-356, 1993
A;Title: Cloning and expression of odorant receptors.
A;Reference number: S29707; MUID:93149273; PMID:7678922
A;Accession: S29710
A;Molecule type: mRNA
A;Residues: 1-307 <RAM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI000004187C C;Superfamily: olfactory receptor OR14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HQDVMKLACDDIRVNVVYGLIVI----ISAIGLDSLLISFSYLLILKTVLGLTREAQAKA 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCL 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAMAF 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENNNITEFILLGLTQDFDGRKALFVIFF---LIYIVTMMGNLLIVVTVIASFSLGSFMY 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GNESSATYFILIGL---PGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMY 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAFG 241
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                                                                                                                                                                                                                                                                                                                                                               TKEIRQRILRLFHVATHA 315
                                                                                                                                                                                                                                                                                                                                                                                                                               LSTCSSHILVVILFFVPCIFMYARPVYNP-----PIDKCITVFYTIITPMLNPLIYTLR 288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FGTCVSHVCAVFIFYVP--FIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVK 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MSPLIVIACTD---TYPIGLTVIANGGVNCIVIFTLLIG-SYGIILRSLKTQSQEGRRKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IRORILEL 308
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CEHMALVQLACGDISINNALGLLTVFLTITADFIFITISYIVILVSIL-RSGKACLKAVN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSY 181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DRYVAICKPLHYLAIMNRRVCITLLIFAWTGGFTHSLIQIVFVYNLPFCGPNVIDHFICD 178
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H.; Beasley, L.; Dionne, V.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         125; Indels
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R;Buck, L.; Axel, R.
Cell 65, 175-187, 1991
A;Title: A novel multigene family may encode odorant receptors: a mole A;Reference number: A23701; MUID:91191556; PMID:1840504
A;Recession: I23701
A;Retatus: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 1-312 <BUC>
A;Ross-references: UNIPROT:P23273; UNIPARC:UP10000041F7F; GB:M64391; C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Proc. Natl. Acad. Sci. U.S.A. 89, 8948-8952, 1992
A,711tle: Spatial pattern of receptor expression in the olfactory epithelium.
A,78cference number: A46247; MUID:93028384; PMID:1384038
A,Accession: A46247
A,Status: preliminary; not compared with conceptual translation
A,78tatus: preliminary; not compared with conceptual translation
A,78cleule type: nucleic acid
A,78csidues: 1-312 < NEFP
A;Cross-references: UNIPROT:P23275; UNIPARC:UPI0000029735; GB:M84005; NID:g200153; PIDN:A,Note: sequence extracted from NCBI backbone (NCBIP:115362)
C;Superfamily: olfactory receptor OR14
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C;Species: Rattus norvegicus (Norway rat)
C;Dats: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LHEPMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTV
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  AICFPLRYTTIMS----TKFCASLVLLLWMLTMTHALLHTL--LIARLSFCEKNVILHFF
                                                                                                                                MLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMBSTVLLAMAFDRYV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VEGRRKAFNTCVSHLVVVFLFYG-----SAIYGYLLPAKSSNQSQGKFISLFYSVVTPMV
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                                          AICHPLRHATVLTLPRVTKIGVAAVV-----RGAALMAPLPVFIKQLPFCRSNILSHSY 181
                                                                                     NLSFSDLCFSSVTMPKLLQNMQSQVPSISYTGCLTQLYFFMVFGDMESFLLVVMAYDRYV
                                                                                                                                                                           NQTLILBFLLLGLPIPSEYHLLFYALFLAMYLTIILGNLLIIVLVRLDSHLHMPMYLFLS
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33.5%;
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34.6%; Pred. No. 1e-30;
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                                                                                                                                                                                                                                                            Pred. No. 3e-30;
4; Mismatches 119; Indels
                                                                                                                                                                                                                                                                              Score 434.5; DB 2;
Pred. No. 3e-30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  312
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                                                                                                                                                                                                                                                                                                      Length 312;
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                                                                                                                                                                                                                                                            Gaps
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    178
                                                                                       124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Jan-1993 #sequence revision 22-Jan-1993
C;Accession: E23701
R;Buck, L.; Axel, R.
Cell 65, 175-187, 1991
A;Title: A novel multigene family may encode odoran
A;Reference number: A23701; MUID:91191556; PMID:184
A;Accession: E23701
olfactory receptor HTPCR92 - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 14-Oct-1997 #sequence_revision 07-Nov-1997 #text_change
C;Accession: JC5624; S58003
R;Vanderhaeghen, P.; Schurmans, S.; Vassart, G.; Parmentier, M.
Biochem. Biophys. Res. Commun. 237, 283-287, 1997
Biochem. Biophys. Res. Commun. 237, 283-287, 1997
A;Title: Molecular cloning and chromosomal mapping of olfactory
A;Reference number: JC5624; MUID:97415789; PMID:9268701
A;Contents: Testis
A;Accession: JC5624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 6
E23701
olfactory receptor I3 - rat
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C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Status: nucleic acid sequence not shown A;Molecule type: mRNA A;Residues: 1-310 <BUC>
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             126 YVAICHPLRHATVLTLPRVTKIGVA---AVVRGAALMAPLPVFIKQLPFCRSNILSHSYC 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 LSNLSFSDLCFSSVTMPKLLQNMRSQDTSIPYGGCLAQTYFFMVFGDMESFLLVAMAYDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLHQDVMKLACDDIRVN-----VVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREA 235
                                                                                                                                                                                                                                                                                                                                             RORILAL 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDR 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NESSATYFILIGLPGLEBAQ--FWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IYKVFSTCGSHLSVVTLFYGTIFGIYLCP--SGNNSTVKEIAMAMMYTVVTPMLNPFIYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QAKAFGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYG
                                                                                                                                                                                                                                                                                                         KRALIRV 302
                                                                                                                                                                                                                                                                                                                                                                                                                                    TCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKEI 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFG 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDISALLKISCSDIYVNELMIYILGGLIIII-----PFLLIVMSYVRIFFSILKFPSIQD
                                                                                                                                                                                                                                                                                                                                                                                            TCGSHLSVVSLFYGTIIGLYLCP-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DLFVLLKLACSDTYINELMIFIMSTLLIIIPFFLIVMSYARIISSILKVPSTQGICKVFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     YVAICFPLHYTSIMS-PKLCTCLVLLLWMLTTSHAMMHTL--LAARLSFCENNVVLNFFC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NOTFITOFLLGLPIPEBHOHLFYALFLV--MYLTTILGNLLIIVLVOLDSOLHTPMYLF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LRNRDMKRALIRV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26.6%; Score 431.5; DB 2; 33.2%; Pred. No. 5.4e-30; ative 71; Mismatches 121;
                                                                                                                                                                                                                                                                                                                                                                                              - AGNNSTVKEMVMAMMYTVVTPMLNPFIYSLRNRDM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ue odorant receptors:
PMID:1840504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121; Indels
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13;

receptor genes expressed

295

177

120

60

þ

molecular

basis for

122

62 65

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C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence_revision 10-Nov
C;Accession: $29708
R;Raming, K.; Krieger, J.; Strotmann, J.; Boel
Nature 361, 353-356, 1993
A;Title: Cloning and expression of odorant re
A;Reference number: $29707; MUID:93149273; PM
A;Accession: $29708
A;Accession: $29708
A;Residues: 1-312 <RAM>
A;Residues: 1-312 <RAM>
A;Cross-references: UNIPARC:UPI0000042028
C;Superfamily: olfactory receptor OR14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Molecule type: mRNA
A;Residues: 1-319 <VANA
A;Residues: 1-319 <VANA
A;Residues: 1-319 <VANA
A;Residues: 1-319 <VANA
A;Cross-references: UNIPROT:Q15622; UNIPARC:UPI0000041DE3; GB:Y10530; NID:g2792017; PID:
A;Vanderhaeghen, P.; Schurmans, S.; Vassart, G.; Parmentier, M.
submitted to the EMBL Data Library, July 1995
A;Description: Male germ cells from several mammalian species express a specific repertor, A;Reference number: S57995
A;Accession: S58003
A;Molecule type: mENA
A;Molecule type: mENA
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F;58-79/Domain: transmembrane #status predicted <TM2>
F;58-79/Domain: transmembrane #status predicted <TM3>
F;101-120/Domain: transmembrane #status predicted <TM3>
F;140-164/Domain: transmembrane #status predicted <TM4>
F;197-218/Domain: transmembrane #status predicted <TM5>
F;237-260/Domain: transmembrane #status predicted <TM5>
F;272-292/Domain: transmembrane #status predicted <TM7>
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A;Residues: 126-292 <VAN>
A;Residues: 126-292 <VAN>
A;Cross-references: UNIPARC:UPI000016B10A; EMBL:X89677; NID:g902337; PIDN:CAA61824.1;
A;Cross-references: UNIPARC:UPI000016B10A; EMBL:X89677; NID:g902337; PIDN:CAA61824.1;
C;Genetics:
A;Map position: 19p13.1
C;Function:
C;Function:
C;Function:
C;Function:
A;Description: involved in control of sperm physiology
C;Superfamily: olfactory receptor OR14
C;Superfamily: olfactory receptor OR14
C;Keywords: G protein-coupled receptor; transmembrane protein
C;Keywords: G protein-coupled receptor or14
C;Keywords: G protein-cou
                                                                                                                                                                                                                                                                                                                                                                                         olfactory receptor OR12 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Sep-1998
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   l Similarity
98; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPV-FIXQLPFCRSNILSHSYC 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DÍKRAL 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FSTCASHLSVVSLFYGAILGVYL--SSAATRNSHSSATASVMYTVVTFMLNPFIYSLRNK 295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTK 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DRFVAICHPL-HYMVIMNPHLCGLLVLASWTMSALYSLLQILMVVRLSFCTALBIPHFFC
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26.5%; Scilarity 31.7%; Pr
Conservative 75;
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Score 429.5; DB 2;
Pred. No. 8.2e-30;
5; Mismatches 119;
                                                                                                                                                                                                                                                                                                                                                     Boekhoff, I.;
                                                                                                                                                                                                                                                                receptors.
PMID:7678922
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      Indels
                                                         Length 312;
                                                                                                                                                                                                                                                                                                                                                        Kubick,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              319;
      17;
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      Gaps
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A;Title: The family of genes encoding odorant receptors A;Reference number: A45774; MUID:93201590; PMID:7916654
A;Accession: G45774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT
G45774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Experimental source: olfactory epithelium A;Note: sequence extracted from NCBI backbone C;Superfamily: olfactory receptor OR14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          odorant receptor 202 - channel catfish
C;Species: Ictalurus punctatus (channel catfish)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-328 <NGA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Ngai, J.; Dowling, M.M.; Buck, L.; Axel, R.; Chess,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62 YIMLMLAASDVLESTITLEKIIARYWEGDGSISEVGCEIQMQEVHYEATVNALVLAVMAE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      64 IFLCMLSGIDILISTSSMPKMLAIFWENSTTIQFDACLLQIFAIHSLSGMESTVLLAMAF 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      66
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Similarity 29.4%; Pred. No. 1.2e-29;
92; Conservative 76; Mismatches 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G45774
                                                            GVKTKEIRQRILR
                                                                                                                     LSTCSSQLIIITLFFLPRCLNYLSSSLGIH-----INADIQILVIMLYSLLPPMINPVIY
                                                                                                                                                                                                                                                                                                                                                                                                              DRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCL 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PEGNITNVKNEVILGPPGLPPNYYGLVSVVMFFVYVCTLIGNCTFFTLFLREKSLQKPMY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PNGNESSATYFILIGLEGLEBAQFWL-AFPLCSLYLIAVLGNLTIIYIVRTEHSLHEEMY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LSNLSFSDLCFSSVTMPKLLQNMQNQDTSITYTGCLTQMYFSMVFAGMEIFLLVSMAYDR
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      CLRTKEAKECLKR
                                                                                                                                                                            FGTCVSHVCAVFIFYVP----FIGLSM-VHRFSKRRDSPLPVILANIYLLVPPVLNPIVY
                                                                                                                                                                                                                                        HVSVTKLACID---RTPYAFPALVSALVMLLTPLAFILFSYGSIIVTVFRTSSTRGRLKT
                                                                                                                                                                                                                                                                                        HQDVMXLACDDIRVNVVYGLIVIISAIGL---DSLLISFSYLLILKTVLGL-TRBAQAKA
                                                                                                                                                                                                                                                                                                                                                        DRYVAVCNPLRYVNIVKESTILGLCVVSWLLAEPTVLTTVIRATSLPYCASNTVIQCYCD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FGTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTK 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALLKLACSDTYINELMIFILGGLLIVI----PFLLIVMTYVQIVCSILKVPSTRAIYKI 237
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RESULT 11
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AJ7286

c) factory receptor I15 - rat
C; Species: Rattus norvegicus (Norway rat)
C; Species: Rattus norvegicus (22-Jan-1993 #text_change 09-Jul-2004
C; Accession: AJ7286
R; Buck, L; Axel, R.
Cell 65, 175-187, 1991
A; Title: A novel multigene family may encode odorant receptors: a molecular basis for ca; Reference number: AZ3701; MUID:91191556; PMID:1840504
A; Reference number: AZ3701 multigene not shown; not compared with conceptual translation A; Residues: 1-314 spuc-
A; Cross-references: UNIPROT:P23274; UNIPARC:UPI0000040A93; GB:M64392; NID:g205845; PIDA C; Keywords: G protein-coupled receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Farmentier, M.; Libert, P.; Schurmans, S.; Schiffmann, S.; Lefort, A.; Eggerickx, D.; Nature 355, 453-455, 1992
A;Title: Expression of members of the putative olfactory receptor gene family in mammal: A;Reference number: $20571; MUID:92131132; PMID:1370859
A;Accession: $20572
A;Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: nucleic acid
A; Residues: 1-314 < PAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             olfactory receptor - human
c;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
C;Accession: $20572
R;Parmentier, M.; Libert, F.; Schurmans, S.; Schiffmann, S.; Lefort, A.; Eg
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Best Local Similarity
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Best Local
                                                                  Matches
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Local Similarity 34.5%; Pred. No. 1.2e-29;
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NESSATYFILIGLPGLEEAQ--FWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYV 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NESSATYFILIGLEGLEBAQFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHEEMYIFLC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AFSTCGSHLSVVSLFYGTVIGLYLC---SSANSSTLKDTVMAMMYTVVTFMLNFFIYSLR 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AFGTCVSHVCAVFIFYVÞFIGLSMVHRFSKRRDSÞL-ÞVILANIYLLVÞÞVLNÞIVYGVK 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AICPPL-HYTAIMSPMLCLALVALSWVLTTFHAMLHTL--LMARLCFCADNVIPHFFCDM 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NQTSISDFLLLGLPIQPEQQNLCYALFLAMYLTTLLGNLLIIVLIRLDSHLHTPMYLFLS
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                                                            26.2%; Score 425.5; DB 2; 33.2%; Pred. No. 1.8e-29; +ive 70; Mismatches 114;
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R;Buck, L.; Axel, R.
Cell 65, 175-187, 1991
A;Title: A novel multigene family may encode odorant receptors: a molecular basis for A;Reference number: A23701; MUID:91191556; PMID:1840504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Raming, K.; Krieger, J.; Strotmann, J.; Boekhoff, I.; Kubick, Nature 361, 353-356, 1993
A;Title: Cloning and expression of odorant receptors.
A;Reference number: S29707; MUID:93149273; PMID:7678922
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence revision 10-Nov-1995 #text_change 31-Dec-2004
C;Accession: S29707; B37286
C;Accession: S29707; B37286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 193-236 <BUC>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-314 < RAM>
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301 IR 302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          126
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                                                                                                                                                                                                                            CLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAF
                                                                                                                                                                                                                                                                                                        YVAICEPLHYMSIMS----PKLCVSLVVLSWVLTTFHAMLHTLLMARLSFCEDNVIPHFF
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                                                         SSCGSHLSVVSLFYGTVIPLYLCP--SSNNSTVKETVMSLMYTLVTPMLNPFIYSLRNRD
                                                                                                                  GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLFVILANIYLLVPFVLNFIVYGVKTKE
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A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-321 <NGA> A;Residues: 1-321 <NGA> A;Cross-references: UNIPROT:09PSJ2; UNIPARC:UPI0000042021 A;Experimental source: olfactory epithelium A;Note: sequence extracted from NCBI backbone (NCBIP:127745) C;Superfamily: olfactory receptor OR14 C;Keywords: olfaction; transmembrane protein
                                                                                                                                                                                       olfactory factor OR37 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Sep-1998
C;Accession: S29711
R;Raming, K.; Krieger, J.; Strotmann, J.; Boekhoff, I.; Kubick, S.; Baumstar
Nature 361, 353-356, 1993
A;Title: Cloning and expression of odorant receptors.
A;Reference number: S29707; MUID:93149273; PMID:7678922
A;Accession: S29711
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C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
C;Accession: H45774
R;Ngai, J.; Dowling, M.M.; Buck, L.; Axel, R.; Chess, A.
Cell 72, 657-666, 1993
A;Title: The family of genes encoding odorant receptors in the channel catfish.
A;Reference number: A45774; MUID:93201590; PMID:7916654
A;Accession: H45774
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C;Species: Ictalurus punctatus (channel catfish)
C;Date: 20-Peb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
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                                                                                                       ;Molecule type: mRNA;Residues: 1-305 <RAM>;Coss-references: UNIFARC:UPI000003FF7B;Cross-references: UNIFARC:UPI000003FF7B;Superfamily: olfactory receptor OR14
                                         Query Match
Best Local Similarity
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                        Conservative
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                                         25.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             320
                        62;
                      Score 418.5; DB 2
Pred. No. 7.2e-29;
2; Mismatches 115
                                                             DB 2;
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                        Gaps
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C(species: Rattus norvegicus (Norway rat)
C(spacies: Rattus norvegicus (Norway rat)
C(space: 10-Sep-1999 #sequence revision 10-Sep-1999 #text_change 09-Jul-2004
C(Accession: S51356; S47014
R(sat, U.; Netrasova, B.; Lancet, D.; Natochin, M.
Rur. J. Biochem. 225, 1157-1168, 1994
RyTitle: Olfactory receptor proteins. Expression, characterization and partial purificati
A; Reference number: S51356; MUID:95045546; PMID:7957207
A. Accession. S61356
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A;Residues: 1-309 <ART;
A;Cross-references: UNIPROT:Q63395; UNIPARC:UPI00000405D8;
R;Gat, U.; Nekrasova, E.; Lancet, D.; Natochin, M.
submitted to the EMBL Data Library, July 1994
A;Description: Olfactory receptor proteins: expression, cha
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A;Accession: S47014
A;Status: preliminary
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A;Status: preliminary
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                                                                                                                                                                                                                                       65 SSLSFVDLCYSTVITPKMLVNFLGKKNFITYSECMAQFFFFAIFVVTEGYLLTVMAYDRY
                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                      AQAKAFGTCVSHVCAVFIFY--VPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPI
                                                                                                                                                                                             VAICHPLRHATVLTLPRVTKIGV------AAVVRGAALMAPLPVFIKQLPFCRSNIL 177
                                                                                                                                                                                                                                                                                 CMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NKDVKAAV 295
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                                                                                                       SHSYCLHQDVMKLACDDIRVNVVYGLIVIISAIG--LDSLLISFSYLLILKTVLGL-TRE
                                                                                                                                                      VAICRPLLYNVIMS-SRICSLLVLVAFSLGLFSAVVHTSAMM------NLSFCKSYII
                                                                                                                                                                                                                                                                                                                         GNHSAVVVFVLVGLTKQPBLLLPLFFLFLVIYVLTVVGNLGMILLIIVSPLLHTPMYYFL
                                                                                                                                                                                                                                                                                                                                                                 GNESSATYFILIGLEGLEEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYIFL
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                                                                  SHYFCDALPLIKLACSNTHLNEL--LIFTIGGLNTLVPTLAVAISYVFIFCSILRIRSSE
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                              25.8%; Score 418.5; DB 1; 33.0%; Pred. No. 7.3e-29;
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Cytogenet. Cell Genet. 63, 200-204, 1993

A;Title: The OLFR1 gene encoding the HGMP07E putative olfactory receptor maps to the A;Reference number: A48413; MUID:93251832; PMID:8097991

A;Accession: A48413

A;Molecule type: DNA

A;Residues: 1-312 <SCH>
A;Cross-references: UNIPROT:P34982; UNIPARC:UPI0000041E06; EMBL:X65857; NID:9425220; IA;Note: sequence extracted from NCBI backbone (NCBIN:131483, NCBIP:131485)

C;Genetics:
                                                                        RESULT 17
S58066
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C;Species: Homo sapiens (man)
C;Date: 31-Dec-1993 #text_change 09-Jul-2004
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: A48413
R;Schurmans, S.; Muscatelli, F.; Miot, F.; Mattei, M.G.; Vassart, G.; Parmentier,
Cytogenet. Cell Genet. 63, 200-204, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   F;101-120/Domain: transmembrane #status predicted <TM3>
F;140-164/Domain: transmembrane #status predicted <TM4>
F;197-219/Domain: transmembrane #status predicted <TM5>
F;197-25/Domain: transmembrane #status predicted <TM5>
F;237-255/Domain: transmembrane #status predicted <TM7>
F;272-291/Domain: transmembrane #status predicted <TM7>
F;5/Binding site: carbohydrate (Asn) (covalent) #status predicted F;137,232,290/Binding site: phosphate (Ser) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: GDB:OLFR1

A;Cross-references: GDB:136274; OMIM:164342

A;Map position: 17p13-17p12

C;Superfamily: olfactory receptor OR14

C;Keywords: G protein-coupled receptor; glycoprotein;

F;26-49/Domain: transmembrane #status predicted <TM1>
F;57-79/Domain: transmembrane #status predicted <TM2>
F;57-79/Domain: transmembrane #status predicted <TM2
    probable olfactory receptor tpcr33 - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1996 #sequence_revision 19-Apr-1996 #t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 FFLANLSFTDLFFVTNTIPKWLVNLQSHNKAISYAGCLTQLYFLVSLVALDNLILAVMAY 120
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97; Conserv
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                                                                                                                                                                  PIYSLRNKDMHGALGRL 303
                                                                                                                                                                                                       IVYGVKTKEIRORILRL 308
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#sequence_revision 19-Apr-1996 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25.5%; Score 413; DB 2; Length 312; 30.6%; Pred. No. 2.2e-28; tive 81; Mismatches 107; Indels
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R.Ngai, J.; Dowling, M.M.; Buck, L.; Axel, R.; Chess, A. Cell 72, 657-666, 1993
A;Title: The family of genes encoding odorant receptors A;Reference number: A45774; MUID:93201590; PMID:7916654
A;Accession: 145774
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A;Residues: 1-160 <VAN>
A;Cross-references: UNIPROT:Q62336; UNIPARC:UPI0000042009; EMBL:X89685; NID:g902668; PID:
C;Superfamily: olfactory receptor OR14
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R;Vanderhaeghen, P.; Schurmans, S.; Vassart, G.; Parmentier, M.
submitted to the EMBL Data Library, July 1995
A;Description: Male germ cells from several mammalian species express a specific repertoi
A;Reference number: S57995
A;Accession: S58066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: UNIPROT:Q9PSJ1; UNIPARC:UPI00000FC739
A;Experimental source: olfactory epithelium
A;Note: sequence extracted from NCBI backbone (NCBIP:1277
C;Superfamily: olfactory receptor OR14
C;Keywords: olfaction; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Ictalurus punctatus (channel catfish)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
C;Accession: 145774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Residues: 1-344 <NGA>
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                                                                                                      181 YCLHQDVMKLACDDIRVN--VVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQ-A
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238 KAFGTCVSHVCAVFIFYVÞFIGLSMVHRFSKRRDSÞLÞVILANIYLLVÞÞVLNÞIVYGVK 297
                                                                                                                                                                                              125 MAYDRVVAICFPLRYNVIVTKBAMTLIIVITWIPSISIIALLVALITRLSFCRSVIINSY 184
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                                                                                                                                                                                                                                                                                                                            65 AKYİTVENLALSÖLGGSSALIPKLIDTELERNQVİSYBACLANMEEVLEEMTVQSLTLLV 124
                                                                                                                                                                                                                                                                                                                                                                                      61 PMYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 ISNPLRYASVLTNNVIIRIGVAITTRATLSLLPLPFPLLKRLNYCPGKILLSHSFCFHADV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 h 25.3%; Score 411; DB 2; Similarity 31.7%; Pred. No. 3.7e-28; 99; Conservative 72; Mismatches 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MKLACADITVNILYGLYVVLSTVGIDSLLIVMSYSLILHTVMGLASPRB-RVRTLNTCVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                      NALSANATFIRPSTFYINGFYNIPHTKYYYAF-LCIAYAVTVLGNSFIMCTIYLARSLHT 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |: || :||:| ||:||:|| | : ::| :||:|| 160
                                                                                                                                                                                                                                                           MAFDRYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NGNESSATY----FILIGLEGGEBAQFWLAFFLCSLYLIAVLGNLTIIYIVRTBHSLHE 60
                                                                FCDHGPILILAGNDKFINRVMAIGCFVVLDCVPF--LLIIVSYICIGIALMNISHGLERR
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50.9%; Pred. No. 1.6e-28;
Live 29; Mismatches 46
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olfactory receptor - dog
C;Species: Canis lupus familiaris (dog)
C;Date: 22-Nov-1993 #sequence_revision 01-Dec-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |Generation | Generation | Gene
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A;Title: Chemoreceptors expressed in taste, A;Reference number: JC5200; MUID:97080538; A;Accession: JC5200
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 27-Feb-1997 #sequence_revision 27-Feb-1997 #text_change
C;Accession: JC5200; PC4302
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A; Residues: 1-311 < THO1>
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,Residues: 146-153;265-272 <THO2>
,Cross-references: UNIPARC:UPI0000178381, UNIPARC:UPI0000178382
,Experimental source: taste bud
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Local Similarity 32.5*; Pred. No. 6.6e-28;
nes 100; Conservative 59; Mismatches 134; Indels 15
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                                                                                                                                                                                                                                                                                                                                                                                                IRORILRL 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MKLSCSDTHVNE-----LVLSGFGGTVLMVPFVSIVISYVHIVFAVLRIQSSGGSSKAFS 237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NOSSVSEFFLRGISGFPEQQQLLYGLFLCMYLVTLTGNVLIILAIGSDPHLHTPMYFFLA 62
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R;Parmentier, M.; Libert, F.; Schurmans, S.; Schiffmann, S.; Lefort, A.; Eggerickx, D.; I Nature 355, 453-455, 1992
A;Title: Expression of members of the putative olfactory receptor gene family in mammaliant A;Reference number: S20571; MUID:92131132; PMID:1370859
A;Accession: S20571; MUID:92131132; PMID:1370859
A;Accession: S20571
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-313 <PAR>
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288
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Walnic B., Godfrey P.A., Buck L.B.;

Walnic B., Godfrey P.A., Buck L.B.;

With human oliactory receptor gene family.";

With human oliactory receptor gene family.";

With human oliactory receptor gene family.";

With human oliactory receptor activity amoutation (TPA) entry.

EMBL/GenBank/DDBJ third party amoutation (TPA) entry.

EMBL/GenBank/DDBJ third party amoutation (TPA) entry.

EMBL/GENBANA/4767.1; -; Genomic DNA.

OR GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004984; F:olfactory receptor activity; IEA.

GO; GO:0004984; F:olfactory receptor activity; IEA.

GO; GO:0004984; F:receptor activity; IEA.

GO; GO:0004984; F:receptor activity; IEA.

GO; GO:000186; F:receptor activity; IEA.

GO; GO:000165; F:receptor activity; IEA.

InterPro; IPR000726; GPCR Rhodpsn.

InterPro; IPR000726; GPCR Rhodpsn.

InterPro; IPR0007276; GPCR Rhodpsn.

PANTHER; PTHR11398; Olfact receptor; 1.

Pfam; PP00001; 7tm l; 1.
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PRINTS; PR00245; OLFACTORYR.

PROSITE; PS00237; G PROTEIN RECEP_F1_1; UNKNOWN_1.

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Olfactory receptor OR11-15.
Homo sapiens (Human).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota, Metazoa; Chordata; Craniata; Primates; Catarrhini; Hominidae;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
 10-OCT-2003 (Rel. 42, Created)
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Q6IF93;
05-JUL-2004 (TrEMBLrel.
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RA Klausner R.D., Collins F.S., Wagner L.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschanko L., Marusina K., Farmer A.A., Rubin G.M., Hsieh F.,

RA Distchenko L., Marusina K., Farmer A.A., Rubin G.M., Hsieh F.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stepleton M.J., Ugdin T.B., Toshiyuki S., Carninci F., Frange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,

RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Willing M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grinwood J., Schmutz J., Myers R.M.,

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RA Rodriguez A.C., Grinwood J., Schmutz J., Myers R.M.,

RA Butterfield Y.S.N., Krzywinski M.I., Skalaka U., Smailus D.E.,

RA Rodriguez A.C., Grinwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grinwood J., Schmutz J., Myers R.M.,

RA RODRELULAR LOCATION: Integral membrane protein.

CC -I- SUNGELLULAR LOCATION: Integral membrane protein.

CC -I- SIMILARITY: Belongs to the G-protein coupled receptor 1 family.

CC -I- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium

CC -I- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium
                                                                               InterPro; IPR000775; GPCR Rhodpsn.

InterPro; IPR000725; Olfact_receptor.

PANTHER; PTHR11398; Olfact_receptor; 1.

Pfam; PP00001; 7tm 1; 1.

PRINTS; PR00237; GPCRRHODOPSN.

PRINTS; PR00237; GPCRRHOTEIN_RECEP_F1_1; 1.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.

G-protein coupled receptor; Glycoprotein; Multigene family; G-protein; Sensory transduction; Transducer; Transmembrane.

TOPO_DOM 1 27 Extracellular (Potential).
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Submitted (JUL-2001) to the EMBL/GenBank/DDBJ
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Tsutsumi S., Aburatani H., Asai K.,
"Genome-wide discovery and analysis
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13-SEP-2005 (Rel. 48, Last annotation updat
Olfactory receptor 51E1.
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BC022401; AAH22401.1; -;
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Best Local &
Matches 316
weng J., Wang J., Cai Y., Ittmann M., Liu M.;

"POGR, a prostate-overexpressed G protein-coupled receptor is sensitive biomarker for human prostate cancers.";

Submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AY775731; AAV54110.1; -; mRNA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004984; F:olfactory receptor activity; IEA.

GO; GO:0004986; P:receptor activity; IEA.

GO; GO:0007186; P:receptor activity; IEA.

R GO; GO:0007186; P:G-protein coupled receptor protein signalin.

R FRINTS; PRO00217; Olfact_receptor.

R PRINTS; PRO00217; Olfact_receptor.
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01-FEB-2005 (TrEMBLrel. 29, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
Prostate overexpressed G protein-coupled receptor.
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Mammalia; Eutheria;
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Q5S4P5;
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Pred. No. 2.6e-113;
1; Mismatches 0;
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Cytoplasmic (Potential).
N-linked (GlCNAc. . .) (Potential).
N-linked (GlCNAc. . .) (Potential).
By similarity.
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Best Local Sim
Matches 316;
               TU-GPCE: a novel putative G protein-coupled receptor over prostate cancer and prostate.";

Biochem. Biophys. Res. Commun. 322:239-249(2004).

RESHL; AXV998056; AAU07956.; Hemo sapiens.

GO; GO:0004962; F:Cintegral to membrane; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0007166; P:G-protein coupled receptor protein signor; GO:0007166; P:G-protein coupled receptor.

GO; GO:0007165; P:G-protein coupled receptor.

GO; GO:0007165; P:G-protein coupled receptor.

RESTITES, FREGUATION PROUPER.

RINTERPORTINE, FTHAI1398; Olfact receptor; 1.

R PAINTS; PRO0237; GFCRRHODOPSN.

R PRINTS; PRO0237; GFCRRHODOPSN.

R PRINTS; PRO0237; GPROTEIN_RECEP_F1_1; 1.

R PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.
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Q66X57;
25-OCT-2004 (
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25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Dresden-G-protein-coupled receptor.
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PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.

Receptor.

SEQUENCE 317 AA; 35271 MW; 8C7293AA7FBCA95C CR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Prostate;

PubMed=15313197; DOI=10.1016/j.bbrc.2004.07.106;

Weigle B., Fuessel S., Ebner R., Temme A., Schmitz M.,

Kiessling A., Rieger M.A., Meye A., Bachmann M., Wirth
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Mammalia; Eutheria;
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01-FBB-2005 (TrEMBLrel. 29, Last annotation update)
01factory receptor MOR18-1 (Olfactory receptor Olfr558)
0verexpressed G protein coupled receptor).
Name=Olfr558,
Mus museculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Bute
                         MEDLINE=22974002; PubMed=14611657; DOI=10.1186/gb-2003-4-11-r71; Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams E.M., Trask B.J.; Walker M., Williams E.M., Trask B.J.; "Odorant receptor expressed sequence tags demonstrate olfactory expression of over 400 genes, extensive alternate splicing and unequal expression levels.";
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NUCLEOTIDE SEQUENCE.
MEDLINE=21864068; PubMed=11875048; DOI=10.1093/hmg/l1.5.535;
Voving J.M., Friedman C., Williams B.M., Ross J.A., Tonnes-Pr
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MEDLINS=21678863; PubMed=11802173;
Zhang X., Firestein S.;
The olfactory receptor gene superfamily
Nat. Neurosci. 5:124-133(2002).
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Matches 295
QSMD65_RAT PRELIMINARY; PRT; 317 AA.

QSMD65;

QSMD65;

TO1-FEB-2005 (TrEMBLrel. 29, Created)

TO1-FEB-2005 (TrEMBLrel. 29, Last sequence update)

TO1-FEB-2005 (TrEMBLrel. 29, Last annotation update)

TO1-FEB-2005 (TrEMBLrel. 29, Last annotation update)

FOOSTATE OVEREXPRESSED G.

FOOSTATE OVEREXPRESSED G.

Rattus norvegicus (Rat).

S Rattus norvegicus (Rat).

S Rattus norvegicus (Rat).

S Rattus norvegicus (Rat).

C Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut

C Eukaryota; Murinae; Rattus.

C Muridae; Murinae; Rattus.
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GO; GO:0004984; F:olfactory receptor activity; IE3
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0007186; P:receptor activity; IEA.
GO; GO:0007165; P:signal transduction; IEA.
InterPro; IPR000276; GPCR Rhodpsn.
InterPro; IPR000725; Olfact receptor.
PANTHER; PTHR11398; Olfact receptor; 1.
PANTHER; PTHR11398; Olfact receptor; 1.
PERINTS; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPCRHODOPSN.
PROSITE; PS00237; GPCRHODOPSN.
PROSITE; PS00237; GPCRHODOPSN.
PROSITE; PS00237; GPCRHODOPSN.
PROSITE; PS00237; GPROTEIN_RECEP_F1_1; UNKNOWN_1
PROSITE; PS00237; GPROTEIN_RECEP_F1_2; 1.
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"POGR, an olfactory G-protein coupled receptor,
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NUCLEOTIDE SEQUENCE.
Sanders K.;
Sanders K.;
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Pred. No. 1.2e-105;
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Length Indels

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Sciurognathi;

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Best Local Sim:
Matches 291;
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
01factory receptor (Fragment).
Gorilla gorilla (gorilla).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Buarchontoglires; Primates; Cat
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                                   MUCLEOTIDE SEQUENCE.
Gilad Y., Wiebe V., Przeworski M., Lancet D., Paabo & "Loss of Olfactory Receptor Genes Coincides with the Full Trichromatic Vision in Primates.";
PLOS Biol. 2:0120-0125(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q6SNB2 9PRIM PRELIMINARY;
                                                                                                                                                                                                                                   NCBI_TaxID=9593;
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ilarity 92.4%; Pred. No. 1.4e-104;
Conservative 11; Mismatches 13;
   AAR19522.1; -; Genomic_DNA
                                                                                                                                                                                                                                                                                                          Chordata; Craniata; Vertebrata; Buteleostomi; Buarchontoglires; Primates; Catarrhini; Hominidae;
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SEQUENCE
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(OBVBV9;

O1-MAR-2002 (TrEMBLrel. 20, Created)

O1-MAR-2002 (TrEMBLrel. 20, Last sequence update)

O1-FEB-2005 (TrEMBLrel. 20, Last sequence update)

O1-FEB-2005 (TrEMBLrel. 20, Last annotation update)

O1-FEB-2005 (TrEMBLrel. 20, Last annotation update)

O1-FEB-2005 (TrEMBLrel. 20, Last annotation update)

O1-FEB-2005 (TrEMBLrel. 20, Last annotation update)

O1-FEB-2005 (TrEMBLrel. 20, Last annotation update)

O1-FEB-2005 (TrEMBLrel. 20, Last sequence)

O1-FEB-2005 (TrEMBLrel. 20, Last sequence)

Sequence (Prostate Prostate                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0004994; F:olfactory receptor act.
GO; GO:0004872; F:receptor act.
GO; GO:0004872; F:receptor act.
GO; GO:0007186; P:G-protein coupled receptor protein signalin..
InterPro; IPR000276; GPCR_Rhodpsn.
InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor;
PANTHER; PRHR11398; Olfact_receptor;
PRINTS; PR00245; OLFACTORYR.
NUCLEOTIDE SEQUENCE Adams M.; Submitted (JAN-2002)
                                                                                    receptor gene
Hum. Mol. Gene
                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
MEDLINE=21676863; PubMed=11802173;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                        Young J.M.,
Trask B.J.;
                                                                                                                                                                                MEDLINE=21864068; PubMed=11875048; DOI=10.1093/hmg/11.5.535;
Young J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Pr
                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                         Nat. Neurosci.
                                                                                                                                                                                                                                                                                         Thang X., Firestein S.;
                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=Olfr78;
                                                                                                                                   "Different evolutionary processes shaped
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                                                                                      Genet. 11:535-546(2002)
  (JAN-2002)
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ilarity 98.7%;
Conservative
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    EMBL/GenBank/DDBJ databases
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Pred. No. 8.6e-76;
1; Mismatches 2;
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RA Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
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RA Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Fleichmann W., Gassterland T., Gissi C., King B., Kochiwa H.,
RA Kuchl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
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RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Gustincich S., Ringwald M., Rodriguez I., Sakamoto N.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
RA Hayashizaki Y.,
RC STRAIN-CSPBL/G; TISSUB-Corebellum, and Skin;
RC STRAIN-CSPBL/G; TISSUB-Corebellum, and Skin;
RX MEDLINE-22354683; PubMed=12466851; DOI=10.0108/nature01266;
RA MEDLINE-22354683; PubMed=12466851; DOI=10.0108/nature01266;
RA MEDLINE-22354683; PubMed=12.010.0108/nature01266;
RA MEDLINE-22354683; PubMed=12.0108/nature01266;
RA Mikaido I., Geato N., Kaito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA Nikaido I., Osato N., Saito R., Nogami A., Schonbach C., Gojobori T.,
RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA Schimil L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,
RA Baldarelli R., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Balke J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Balda E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S.,
RA Gasaterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,
RA Gasaterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,
RA Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Kanai A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
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RA Magashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
RA Ravasi T., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,
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Submitted
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STRAIN=C57BL/6J; TISSUE=Cerebellum, and Skin;
MEDLINE=21085660; PubMed=11217851; DOI=10.1038/35055500;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=C57BL/6J; TISSUE=Cerebellum, and Skin;
MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
Carninci P., Hayashizaki Y.;
"High-efficiency full-length cDNA cloning.";
Meth. Enzymol. 303:19-44(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        species."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human olfactory receptor that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=21564169; PubMed=11707321; DOI=10.1016/S0378-1119(01)00709-0; Yuan T.T., Toy P., McClary J.A., Lin R.J., Miyamoto N.G., Kretschmer P.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ional annotation of a 409:685-690(2001).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ដ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      full-length mouse cDNA collection.";
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RC STRAIN-CSTBLIGG; TISSUB=Cerebellum, and Skin;
RC STRAIN-CSTBLIGG; TISSUB=Cerebellum, and Skin;
RA Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P.,
RA Pukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,
RA Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,
RA Hori F., Imotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,
RA Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S.,
RA Kurihara C., Matsuyama T., Miyazaki R., Konno H., Sano H.,
RA Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,
RA Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,
RA Sasaki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M.,
RA Tagawa A., Takahashi F., Takaku-Akahira S., Takeda Y., Tanaka T.,
RA Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.;
RA Tomaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y.;
RA Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.

DR EMBL; AY073011; AAL60674.1; -; Genomic DNA.

DR EMBL; AY073011; AAL60674.1; -; Genomic DNA.

DR EMBL; AY073011; AAL60674.1; -; Genomic DNA.

DR EMBL; AY073012; AAL75109.1; -; mRNA.

EMBL; AY073013; AAL75109.1; -; mRNA.

EMBL; AX036356; BAC29396.1; -; mRNA.

EMBL; AX036356; BAC29396.1; -; mRNA.

DR EMBL; AX036356; BAC29396.1; -; mRNA.

EMBL; AX036356; BAC29396.1; -; mRNA.

DR EMBL; AX036350; Olfact receptor.

DR GO; GO:0016021; C:integral to membrane; TAS.

DR FATM; PF00001; 7tm 1; 1.

DR PATTER; PTHR11398; Olfact receptor: 1.

DR PATTER; PTHR11398; Olfact receptor: 1.

DR PATTER; PTHR11398; Olfact receptor: 1.
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Best Local
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MEDLINB=20530913; PubMed=11076861; DOI=10.1101/gr.152600;
MEDLINB=20530913; PubMed=11076861; DOI=10.1101/gr.152600;
Shibata K., Itch M., Alzawa K., Nagaoka S., Sasaki N., Carninci P.,
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Youno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itch M.,
Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,
Pujiwake S., Inoue K., Togawa Y., Izawa M., Ohara B., Watchiki M.,
Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,
Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;
"RIKEN integrated sequence analysis (RISA) system-384-format
sequencing pipeline with 384 multicapillary sequencer.";
Genome Res. 10:1757-1771(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=C57BL/6J; TISSUB=Cerebellum, and Skin; MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100; Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K., Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.; Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.; Proparatization and subtraction of cap-trapper-selected cDNAs prepare full-length cDNA libraries for rapid discovery of new Genome Res. 10:1617-1630(2000).
                                                                                           PRINTS; PRO0237; GEPCRHODOPSN.
PRINTS; PR00245; OLFACTORYR.
PROSITE; PS00237; G PROTEIN RECEP_F1_1;
PROSITE; PS50282; G_PROTEIN_RECEP_F1_2;
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Similarity
                                                             320
                                                           Ŗ,
                                                             35578 MW;
60.7%;
59.8%;
Score 985;
Pred. No.
                                                           7161ACA8F4328959 CRC64;
    5e-66;
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                   DB 2;
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                       Length 320;
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OSIB2 HUMAN STANDARD,
Q9H255;
Q9H255;
Q9H255;
16-OCT-2001 (Rel. 40, Last sequence update)
13-SEP-2001 (Rel. 48, Last annotation update)
13-SEP-2005 (Rel. 48, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
Xia C., Ma W., Liu M.;
"Identification of a prostate-specific G-protein coupled receptor (PSGR) that interacts with G alpha subunit in yeast two hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Xu L.L., Stackhouse B.G., Florence K., Zhang W., Shanmugam N., Sesterhenn I.A., Zou Z., Srikantan V., Augustus M., Roschke V., Scarter K., McLeod D.G., Mouli J.W., Soppet D., Srivastava S.; "PSGR, a novel prostate-specific gene with homology to a G protein-coupled receptor, is overexpressed in prostate cancer."; Cancer Res. 60:6568-6572(2000).
                                                                                                                                                                                                                      MEDLINE=21564169; PubMed=11707321; DOI=10.1016/S0378-1119(01)00709-0; Yuan T.T., Toy P., McClary J.A., Lin R.J., Miyamoto N.G., Kretsechmer P.J.; Kretsechmer P.J.; Characterization of an evolutionarily conserved "Cloning and genetic characterization of an evolutionarily conserved human olfactory receptor that is differentially expressed across
  MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PSGR) that assay.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
MEDLINE=21002538; PubMed=11118034;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
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Homo sapiens (Human).
                                                                                                                 NUCLEOTIDE SEQUENCE [LARGE
                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (APR-2001) to
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                                                                                                                 SCALE MRNA]
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  Schuler G.D.
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RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsich F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Rownstein M.J., McEwan R.J., Abramson R.D., Mullahy S.J.,
RASS S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RASS S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
VIllalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
VILLAION D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
VILLAION D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
VA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
VA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
VA Blakesley R.W., Touchman J.W., Green B.D., Dickeon M.C.,
VA Butterfield Y.S.N., Xrzywinski M.I., Skahska U., Smallus D.E.,
VG Generation and initial analysis of more than 15,000 full-length human
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PRINTS; PR00245; OLFACTORYR.

PROSITE; PS00237; G PROTEIN RECEP_F1_2; 1.

PROSITE; PS50262; G PROTEIN_RECEP_F1_2; 1.

G-protein coupled receptor; Glycoprotein; Multigene family; Receptor; Sensory transduction; Transducer; Transmembrane.

TOPO_DOM 1 24 Extracellular (Potential).
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EMBL;
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GO; GO:0004984; F:olfactory receptor activity; NAS.
GO; GO:0007608; P:perception of smell; NAS.
InterPro; IPR000276; GPCR Rhodpsn.
InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                    H-InvDB; HIX0009385; -
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AF369708; AAK38728.1; -; mRNA.
AY333942; AAK37550.1; -; mRNA.
BC020768; AAH20768.1; -; mRNA.
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Cytoplasmic (Potential).

2 (Potential).

Extracellular (Potential).

3 (Potential).

Cytoplasmic (Potential).

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                                                    Extracellular (Potential).
5 (Potential).
Cytoplasmic (Potential).
6 (Potential).
Extracellular (Potential).
7 (Potential).
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Query Match Best Local

Similarity

60.4**%**;

Score Pred.

979; DB 1; No. 1.4e-65;

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IT "The human olfactory receptor gene family.";

IT Proc. Natl. Acad. Sci. U.S.A. 101:2584-2589 (2004).

IN EMBL; BK004368; DAA04766.1; -; Genomic_DNA.

R GO; GO:00016021; C:integral to membrane; IEA.

RGO; GO:0004984; F:olfactory receptor activity; IEA.

RGO; GO:0004987; F:receptor activity; IEA.

RGO; GO:0007186; P:G-protein coupled receptor protein signalin. .; IEA.

RGO; GO:0007186; P:G-protein coupled receptor protein signalin. .; IEA.

RGO; GO:0007187; F:GCRRhodpsn.

RINTERPROJUZIT; GFCRRHODOPSN.

PRINTS; PR00237; GFCRRHODOPSN.

PRINTS; PR00237; GFCRRHODOPSN.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; UNKNOWN_1.

PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.
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Best Local S
Matches 182
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
01factory receptor OR11-16.
Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Metazoa; Chordata; Craniata; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
PubMed=14983052; DOI=10.1073/pnas.0307882100;
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Q61F94;
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                                                                                                                                                                                                                                                                                              Score 979; DB 2; Length 320; Pred. No. 1.4e-65; Mismatches 67; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Raming K., Konzelmann S., Breer H.;
"Identification of a movel G-protein coupled receptor expressed distinct brain regions and a defined olfactory zone.";
Recept. Channels 6:141-151(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rattus norvegicus (Rat).
Ruttus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Muroidea; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
Olfactory receptor 51E2 (G-protein coupled receptor RAIc)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AF079864; AAD12761.1; -;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FUNCTION: Putative odorant receptor.
SUBCELLULAR LOCATION: Integral membrane protein.
TISSUE SPECIFICITY: Expressed only in some areas
in the olfactory epithelium.
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NUCLEOTIDE SEQUENCE.
MEDLINE=21676863; PubMed=11802173;
Zhang X., Firestein S.;
"The olfactory receptor gene superfamily Nat. Neurosci. 5:124-133(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OBVF09 MOUSE PRELIMINARY; PRT; 322 AA.

(BVF09)

01-MAR-2002 (TrEMBLrel. 20, Created)

01-MAR-2002 (TrEMBLrel. 20, Last sequence update)

01-PEB-2005 (TrEMBLrel. 29, Last annotation update)

01factory receptor MOR18-3 (Olfactory receptor Olfr557).
                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
MEDLINE-21864068; PubMed-11875048; DOI=10.1093/hmg/11.5.535;
Young J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Pr
Trask B.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
MCBI_TaxID=10090;
NUCLEOTIDE SEQUENCE.
MEDLINE=22974002; PubMed=14611657; DOI=10.1186/gb-2003-4-11-r71;
                                                                                                                                                    receptor gene fan
Hum. Mol. Genet.
                                                                                                                                                                         "Different evolutionary processes receptor gene families.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=Olfr557;
Mus musculus (Mouse).
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Extracellular (Potential).
7 (Potential).
7 (Potential).
Cytoplasmic (Potential).
N-linked (GlONAC. . .) (Potential).
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Pred. No. 2e-65;
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Best Local :
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InterPro; IPR000725; Olfact_receptor.
PANTHER, PTHR11398; Olfact_receptor; 1.
Pfam; PF00001; 7tm_1; 1.
PRINTS; PR00237; GFCRRHODDESN.
PRINTS; PR00237; GFCRTORYR.
PROSITE; PR00237; G PROTEIN_RECEP_F1_1; UN
PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.
                                                                                 OSID1 HUMAN STANDARD; PRT; 324 AA. (9NGF3; 13-SEP-2005 (Rel. 48, Created)
13-SEP-2005 (Rel. 48, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
01factory receptor SID1 (Olfactory receptor OR11-14).
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GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004984; F:olfactory receptor activity; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0007186; P:G-protein coupled receptor protein

GO; GO:0007186; P:signal transduction; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams E.M., Trask B.J., Walker M., Williams E.M., Trask B.J., Good and traceptor expressed sequence tags demonstrate olfactory expression of over 400 genes, extensive alternate splicing and unequal expression levels."
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EMBL; AY073732; AAL61395.1; -; Genomic DNA.
Ensembl; ENSMUSG00000046017; Mus musculus.
                                               Bukaryota; Metazoa;
                                                                        Name=OR51D1;
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 NCBI_TaxID=9606;
                               Mammalia; Eutheria;
                                                         Homo sapiens (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                            KBIRSRVIRME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     322 AA; 35612 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                             Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                            317
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9; Mismatches
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Mainic B., Godfrey P.A., Buck L.B.;
"The human olfactory receptor gene family.";
Proc. Natl. Acad. Sci. U.S.A. 101:2584-2589(2004).
-!- FUNCTION: Putative odorant receptor.
-!- SUNCELULAR LOCATION: Integral membrane protein.
-!- SUNCELULAR LOCATION: Integral membrane protein.
-!- SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
-!- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium
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InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor; 1.
Pfam; PP00001; 7tm 1; 1.
PRINTS; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPCRRHODOPSN.
PRINTS; PR00237; GPROTEIN_RECEP_F1_1; 1.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_2; 1.
G-protein_coupled_receptor; Multigene_family; Olfaction; Receptor;
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between
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Tsutsumi S., Aburatani H.,
"Genome-wide discovery and
                                                                                                                                                                                             TOPO DOM
                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; AB065855; BAC06073.1; -;
EMBL; BK004370; DAA04768.1; -;
Ensembl; ENSG00000197428; Homo
HGNC; HGNC:15193; OR51D1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases
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as long a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WWW="http://bip.weizmann.ac.il/cgi-bin/HORDE/showGene.pl?key=symbol&value=ORS
                    125
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                                                  FLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFD 124
                                                                                             NGNESSATYFILIGLPGL-EEAQFWLAFPLCSLYLIAVLGNLTIIYIVRTEHSLHEPMYI
                    RYVAICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLH 184
                                       FLAMLSTIDLVLSSITMFKMASLFLMGIQEIEFNICLAQMFLIHALSAVESAVLLAMAFD
                                                                               NGNLVHAAYFLLVGIPGLGPTIHFWLAFPLCFMYALATLGNLTIVLIIRVERRLHEPMYL 74
 SEQUENCE.

jato T., Okouchi I., Arita M., Futami jato T., Okouchi I., Arita M., Futami j., Aburatani H., Asai K., Akiyama Y.;

j., Aburatani H., Asai K., Akiyama selde discovery and analysis of human se
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1 (Potential).

Cytoplasmic (Potential).

2 (Potential).

Extracellular (Potential).

3 (Potential).

Cytoplasmic (Potential).

4 (Potential).

Extracellular (Potential).

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7 (Potential).

Extracellular (Potential).

7 (Potential).
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Pred. No. 1.9e
52; Mismatches
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By similarity.
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Trask B.J.;
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EMBL; AY07283; AAL66646.1; -; Genomic DNA.
EMBL; AY317668; AAP71047.1; -; Genomic DNA.
Ensembl; ENSMUSG00000049196; Mus musculus.
MGI; MGI:3030389; Olfr555.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:004984; F:olfactory receptor activity; IEA.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0007186; P:G-protein coupled receptor prot.
GO; GO:0007165; P:signal transduction; IEA.
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01-MAR-2002 (TrEMBLrel. 20, L
01-FEB-2005 (TrEMBLrel. 29, L
Olfactory receptor MOR10-1 (O
Name=Olfr555;
Mus musculus (Mouse).
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MEDLINE=21676863; PubMed=11802173;
Zhang X., Firestein S.;
The olfactory receptor gene super
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Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
                                                  InterPro; IPR000276; GPCR Rhodpsn.
InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                       expression of over 400 genes, expression levels.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Young J.M., Shykind B.M., Lane R.P., Walker M., Williams E.M., Trask B.J.; "Odorant receptor expressed sequence
                                                                                                                                                                                                                                                                                                                    Submitted (JUN-2003) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE
MEDLINE=22974002; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor gene families.
Hum. Mol. Genet. 11:535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Different evolutionary processes shaped the mouse and human olfactory
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MEDLINE=21864068; Pul
Young J.M., Friedman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=10090;
                                       PF00001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                       4:R71-R71(2003).
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; 7tm l; 1.
37; GPCRRHODOPSN.
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man C., Williams E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=14611657;
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Euarchontoglires; Glires; Rodentia; Sciurogna
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Last annotation update)
(Olfactory receptor Olfr555).
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., Tonnes-Priddy L., Ross J.A.,
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Best Local S
Matches 168
                                                                                                                                                                                                                                                                                                                                                                                OBVH11;
01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
Olfactory receptor MOR7-2 (Olfactory receptor Olfr577).
NUCLEOTIDE SEQUENCE.

MEDLINB-22974002; pubMed=14611657; DOI=10.1186/gb-2003-4-11-r71; Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A., Walker M., Williams E.M., Trask B.J.;

"Odorant receptor expressed form:"
                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.
MEDIJNS=21676863; PubMed=11802173;
Zhang X., Firestein S.;
"The olfactory receptor gene superfamily Nat. Neurosci. 5:124-133(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q8VH11_MOUSE PRELIMINARY;
                                                                                                                                                                                                                                                                                                                        Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognathi;
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                Name=Olfr577;
Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRINTS; PR00245; OLFACTORYR.

PROSITE; PS00237; G PROTEIN RECEP F1 1;
PROSITE; PS50262; G PROTEIN RECEP F1 2;
                                                                                                                              receptor gene families.";
Hum. Mol. Genet. 11:535-546(2002)
                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
MEDLINE=2164068; PubMed=11875048; DOI=10.1093/hmg/l1.5.535;
Young J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Pr
Trask B.J.;
                                                                                                     NUCLEOTIDE SEQUENCE
                                                                                                                                                     "Different evolutionary processes
                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10090;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOUSE
15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 914.5; DB 2; Length 315; Pred. No. 9.7e-61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                        shaped the
                                                                                                                                                                                                                                                                                                                                                                                                                                                         312
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                                                                                                                                                                                                                                                of the mouse.";
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                                                                                                                                                         mouse
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                                                                                                                                                         human olfactory
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RASULT 16
OS1G2 HUMA
ID OS1G2
AC OBNGK
DT 10-OC
DT 10-SE
DE Olfac
GN Name=
OS Homo
OC BUKAR
OC HOmo
OC MAMMA
OC HOMO
OC NCBI
RN ULL
RP NUCLE
RA SLWA
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Best Local S
Matches 165
                                                                                                                                                                                                                                                                              O51G2 HUMAN
Q8NGKO;
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Receptor.
SEQUENCE
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PRINTS; PR00237; GPCRHODOPSN.

PRINTS; PR00245; OLFACTOXYR.

PROSITE; PS00237; G PROTEIN RECEP_F1_1;

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GO; GO:001021; C:integral to membrane; IEA.
GO; GO:001021; C:integral to membrane; IEA.
GO; GO:0004984; F:olfactory receptor activity;
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0007186; P:G-protein coupled receptor px
GO; GO:0007165; P:signal transduction; IEA.
InterPro; IPR000276; GPCR Rhodpsn.
InterPro; IPR000725; Olfact_receptor.
PANTHER; PTHR11398; Olfact_receptor; 1.
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GO; GO:0016021; C:integral
GO; GO:0004984; F:olfactor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (JUN-2003) to the EMBL/GenBank/DDBJ EMBL; AY072977; AAL60640.1; -; Genomic_DNA. EMBL; AY317692; AAP71065.1; -; Genomic_DNA. Ensembl; ENSMUSG00000043354; Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression of over 400 genes, extensive alternate splicing and unequal expression levels.";
Genome Biol. 4:R71-R71(2003).
[5]
                                                                                                                                                                                     Olfactory receptor Name=OR51G2;
                                                                                                                    Eukaryota; Metazoa;
Mammalia; Butheria;
                                                                                                                                                                                                        10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last seq)
13-SEP-2005 (Rel. 48, Last ann)
Olfactory receptor 51G2.
NUCLEOTIDE SEQUENCE.
Suwa M., Sato T., Ok
                                                                  NCBI_TaxID=9606
                                                                                                                                                                 omo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                           NAMOH
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                                                                                                                                                                                                                                                                                                                                    STANDARD;
  Okouchi
                                                                                                                                                                                                                                                                                                                                                                                                                                                       310
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                                                                                                                    Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                    sequence update)
annotation update)
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PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.

G-protein coupled receptor; Glycoprotein; Multigene family; Olfaction; Receptor; Sensory transduction; Transducer; Transmembrane.

Receptor; Sensory transduction; Transducer; Obtential).
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InterPro; IPRO00276; GPCR Rhodpsn.
InterPro; IPRO00275; Olfact receptor.
PANTHER; PTHR11398; Olfact receptor;
Pfam; PF00001; 7tm 1; 1.
PRINTS; PR00237; GFCRHODDPSN.
PRINTS; PR00237; OLFACTORYR.
PRINTS; PR00245; OLFACTORYR.
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-i- SUBCELULAR LOCATION: Integral membrane protein.
-i- SIMILARITY: Belongs to the G-protein coupled receptor 1 family.
-i- DATABASE: NAME-Human Olfactory Receptor Data Exploratorium
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
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"Genome-wide discovery and analysis of human seven transmembrane helix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     as long as its content is in no way modified and this statement
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Extracellular (Potential).

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Cytoplasmic (Potential).
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3; Mismatches 81
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ID Q8VH17 MOUSE PRELIMINARY;
AC Q8VH17;
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"The human olfactory receptor gene family.";

Proc. Natl. Acad. Sci. U.S.A. 101:2584-2589 (2004).

EMBL; BK004285; DAA04683.1; -; Genomic DNA.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0004994; F:olfactory receptor activity; IEA.

GO; GO:0004992; F:receptor activity; IEA.

GO; GO:0004916; P:olfactory receptor protein |

InterPro; IPR000276; GPCR Rhodgan.

InterPro; IPR000725; Olfact_receptor.

PRINTS; PR00237; GFCRRHODOPSN.

PRINTS; PR00237; GFCRRHODOPSN.

PRINTS; PR00237; GFCRRHODOPSN.
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SEQUENCE
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Q61FH7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PubMed=14983052; DOI=10.1073/pnas.0307882100; Malnic B., Godfrey P.A., Buck L.B.;
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Submitted (JUN-2003) to the EMBL/GenBank/DDBJ datal
EMBL; AY072971; AAL60634.1; -; Genomic DNA.

R EMBL; AY317693; AAP71066.1; -; Genomic DNA.

R GO; GO:00014021; C:integral to membrane; IEA.

R GO; GO:0004984; F:olfactory receptor Extra Prote GO; GO:0004872; F:receptor activity; IEA.

R GO; GO:0004872; F:receptor activity; IEA.

R GO; GO:0004872; F:receptor activity; IEA.

R GO; GO:0007186; P:signal transduction; IEA.

R GO; GO:0007165; P:signal transduction; IEA.

R GO; GO:
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MEDLINE=21676863; Pubmed=11802173;
Zhang X., Firestein S.;
"The olfactory receptor gene superfamily Nat. Neurosci. 5:124-133(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
01factory receptor MOR7-1 (Olfactory receptor Olfr5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                     LCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDR
                                                                    LAMLALTDLGLCLSTLPTVLGIFWFDARBIGIPACFTQLFFIHTLSLVBSSVLLSMSFDR
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AC Q8VGZE,
DT 01-MAR-2002 (TREMBLIFEL 20, LE
DT 01-MAR-2002 (TREMBLIFEL 20, LE
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DT 01-FEB-2005 (TREMBLIFEL 20, LE
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OX MUCLEOTIDE SEQUENCE.

RY MEDLINE-21664063; PubMed-1180;
RX MEDLINE-21664068; PubMed-1187;
RX MEDLINE-2164068; PubMed-1461;
RX MEDLINE-22974002; to the E
RI "Different evolutionary proces
RT "CIFCOTIDE SEQUENCE.
RA Adams M.;
RI Submitted (JAN-2002) to the E
RX MEDLINE-22974002; PubMed-1461;
RX MEDLINE-22974002; PubMed-1461;
RX MEDLINE-22974002; PubMed-1461;
RX MEDLINE-22974002; PubMed-1461;
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RX MEDLINE-22974002; PubMed-1461;
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     Submitted (JUN-2003) to the EMBL/GenBank/DDBJ database EMBL; AY0772994; AAL60657.1; -; Genomic DNA. EMBL; AY317676; AAP71055.1; -; Genomic DNA. Ensembl; ENSMUSG0000051711; Mus musculus. MGI; MGI:3030395; Olfr561.

GO; GO:0016021; C:integral to membrane; IEA. GO; GO:0004984; F:olfactory receptor activity; IEA. GO; GO:0004982; F:receptor activity; IEA. GO; GO:0007186; P:G-protein coupled receptor protein s GO; GO:0007186; P:signal transduction; IEA. InterPro; IPR000725; Olfact receptor. PANTHER; PTHR11398; Olfact receptor; 1.
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Mus musculus (Mouse).
Mus musculus (Mouse).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalla; Butheria; Buarchontoglires; Glires; Rodentia; Sciurognat
Muridae; Murinae; Mus.
MCBI TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIINE=22974002; PubMed=14611657; DOI=10.1186/gb-2003-4-11-r71 Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A. Walker M., Williams E.M., Trask B.J.; Walker M., Williams E.M., Trask B.J.; Gemonstrate olfactory "Odorant receptor expressed sequence tags demonstrate olfactory expression of over 400 genes, extensive alternate splicing and 1
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MEDLINE=21676863; PubMed=11802173;
Zhang X., Firestein S.;
The olfactory receptor gene super
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MEDLINE=21864068; PubMed=11875048;
MEDLINE=21864068; PubMed=11875048;
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(Olfactory receptor Ol
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., Tonnes-Priddy L., Ross J.A.
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01-MAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-PEB-2005 (TrEMBLrel. 29, Last annotation update)
Olfactory receptor MOR30-1 (Olfactory receptor Olfr
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PRINTS; PR00237; GFCRRHODOPSN.

PRINTS; PR00245; OLFACTORYR.

PROSITE; PS00237; G PROTEIN RECEP_F1_1;

PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;

Receptor.
                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE.
MEDLINE=21676863; PubMed=11802173;
Zhang X., Firestein S.;
"The olfactory receptor gene super
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Z2 MOUSE
QBVGZ2_MOUSE PRELIMINARY;
                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
MEDLINE=21864068, PubMed=11875048, DOI=10.1093/hmg/11.5.535;
Young J.M., Friedman C., Williams E.M., Ross J.A., Tonnes-Priddy L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
  MEDLINE=22974002; PubMed=14611657; DOI=10.1186/gb-2003-4-11-r71; Young J.M., Shykind B.M., Lane R.P., Tonnes-Priddy L., Ross J.A.
                                                                                         Submitted (JAN-2002)
                                                                                                                                                                                             "Different evolutionary processes receptor gene families.";
                                                                                                                                                                                                                                             Young J.M.,
Trask B.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Name=Olfr569;
                                             NUCLEOTIDE SEQUENCE
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162; Conserv
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                                                                                                                                    SEQUENCE.
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Matches 155; Conservative
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GO; GO:0016021; C:integral to membrane; IRA.
GO; GO:0004984; F:olfactory receptor activity; IE.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0007186; P:G-protein coupled receptor prot.
GO; GO:0007165; P:signal transduction; IEA.
Interpro; IPR000276; GPCR Rhodgen.
Interpro; IPR000276; GPCR Rhodgen.
Interpro; IPR000725; Olfact receptor.
PANTHER; PTHR11398; Olfact receptor; 1.
Pfam; PF00001; 7tm 1; 1.
                                                                                                                                                                                                                                                                                                                                                     Receptor.
SEQUENCE
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PRINTS; PR00245; OLFACTORYR.

PROSITE; PS00237; G PROTEIN RECEP_F1_1;

PROSITE; PS50262; G PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (JUN-2003) to the EMBL/GenBank/DDBJ EMBL; AY072998; AAL60661.1; -; Genomic_DNA. EMBL; AY317684; AAP71059.1; -; Genomic_DNA. Ensembl; ENSMUSG0000062142; Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Walker M., Williams E.M., Tras
"Odorant receptor expressed ss
expression of over 400 genes,
expression levels.";
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50.2%; Pred. No. 1.2e-57;
tive 68; Mismatches 85
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'cgn2_6/ptodata/1/iaa/5_COMB.pep:*

'cgn2_6/ptodata/1/iaa/6_COMB.pep:*

'cgn2_6/ptodata/1/iaa/H_COMB.pep:*

'cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

'cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

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ALIGNMENTS

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APPLICANT: Wang, Aljun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THI
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
TYPE: PRT
RESULT 2
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; Sequence 920, Application US/10012896
; Patent No. 6943236
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Best Local :
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Vedvick, Thomas S.
Carter, Darrick
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Kalos, Michael D.
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; ORGANISM: Homo sapiens
US-10-012-896-920
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 1011
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 920
LENGTH: 318
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Best Local Similarity 99.7%;
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APPLICANT:
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APPLICANT: Dillon, Davin C.
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                                                                                                                                             GTCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLFVILANIYLLVPFVLNPIVYGVKTKE 300
                                                                                                                                                                                                     YCLHQDVMKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGLTREAQAKAF
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                                                                              IRQRILRLFHVATHASEP 318
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                                                             IRQRILRLFHVATHASEP
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Day, Craig H.
Vedvick, Thomas (
Carter, Darrick
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Wantanabe, Yoshihiro
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McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
Henderson, Robert A.
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Kalos, Michael D.
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Stolk, John A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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Pred. No. 1e-139;
1; Mismatches
                                                             318
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Sequence 2, Application Patent No. 6790631 GENERAL INFORMATION: APPLICANT: Arthur B. F

US/09680728

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GENERAL INFORMATION:

APPLICANT: Arthur B. Raitano

APPLICANT: Apaniel E.H. Afar

APPLICANT: Aya Jakobovits

APPLICANT: Aya Jakobovits

APPLICANT: Mary Faris

APPLICANT: Steve Chappell Mitchell

APPLICANT: Steve Chappell Mitchell

APPLICANT: Douglas C. Saffran

TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR

TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND

FILE REFERENCE: 511582002410

CURRENT APPLICATION NUMBER: US/10/017,066A

CURRENT PILING DATE: 2002-05-28

PRIOR APPLICATION NUMBER: US 09/680,728

PRIOR APPLICATION NUMBER: 60/157,902

PRIOR APPLICATION NUMBER: 60/157,902

PRIOR APPLICATION NUMBER: 60/157,902

PRIOR APPLICATION NUMBER: 60/157,902
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; ORGANISM: Homo Sapiens
US-09-680-728-2
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10017066A Patent No. 6838258
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Best Local
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LENGTH: 317
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APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
FILE REFERENCE: 129.24USUI
CURRENT APPLICATION NUMBER: US/09/680,728
CURRENT FILING DATE: 2000-10-05
FRIOR APPLICATION NUMBER: 60/157,902
PRIOR PILING DATE: 1999-10-05
NUMBER OF SEQ ID NOS: 50
NUMBER OF SEQ ID NOS: 50
NUMBER OF SEQ ID NOS: 50
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APPLICANT:
APPLICANT:
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 MYIFLCMLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TCVSHVCAVFIFYVPFIGLSMVHRFSKRRDSPLPVILANIYLLVPPVLNPIVYGVKTKEI 301
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Mary Faris
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang Yuqui
APPLICANT: Reed, Steven G.
APPLICANT: Ready Michael
APPLICANT: Ralos, Michael
APPLICANT: Retter, Mark
APPLICANT: Solk, John
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILL REFERENCE: 210121.427C9
CURRENT APPLICATION MUMDER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
                                                                                                                                                                                         ; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 527
; LENGTH: 320
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-439-313-527
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US-09-439-313-527
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                                                                                                             Query Match 60.4%; Score 979; DB 2; Length 320; Best Local Similarity 59.5%; Pred. No. 2.1e-81; Matches 182; Conservative 55; Mismatches 67; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 527,
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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TYPE: PRT
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             301
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68 MLSGIDILISTSSMPKMLAIFWFNSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYV 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5
                                                                           8 NESSATYFILIGLEGLEGAGFWLAFFLCSLYLIAVLGNLTIIYIVRTEHSLHEFMYIFLC
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                                       NFTHAT-FVLIGIPGLEKAHFWVGFPLLSMYVVAMFGNCIVVFIVRTERSLHAPMYLFLC
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FILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND FILE REFERENCE: 210121.42717C17

CURRENT APPLICATION NUMBER: US/99/636,215

CURRENT FILING DATE: 2000-08-10

NUMBER OF SEQ ID NOS: 852

SOFTWARE: Featere for Windows "
SEQ ID NO S27

LENGTH: 320

Type: PPT
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                                                                                                                                                                                                                                                                   ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-636-215-527
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US-09-636-215-527
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Best Local S
Matches 182
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60.4%; Score 979; DB 2; Length 320;
Local Similarity 59.5%; Pred. No. 2.1e-81;
hes 182; Conservative 55; Mismatches 67; Indels
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Dillon, Davin C.
Mitcham, Jennifer L.
Harlocker, Susan L.
MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSH
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
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US-09-685-166A-527
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US-09-685-166A-527
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APPLICANT: Millon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Hatlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Kalos, Michael D.
APPLICANT: Kalos, Michael D.
APPLICANT: Renger, Gary R.
APPLICANT: Retter, Marc W.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Wedvick, Thomas S.
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Li, Samuel
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Best Local :
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Similarity 59.5%; Pred. No. 2.1e-81;
82; Conservative 55; Mismatches 67
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US-09-679-426-527

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APPLICANT: Wang, Aijun
APPLICANT: Skekky, Yasir A.W.
APPLICANT: Skekky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C20
CURRENT APPLICATION NUMBER: US/09/679,426
CURRENT APPLICATION NUMBER: US/09/679,426
CURRENT FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 895
SOFTWARE: PSEC SOFTWARE: COMPOSITION OF SEQ ID NO 527
LENGTH: 320
TYPE: PRT
ORGANISM: Homo sapiens
US-09-679-426-527
RESULT 9
US-09-759-143-527
J Sequence 527, Application US/09759143
Patent No. 6800746
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
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Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
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Carter, Darrick
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Mitcham, Jennifer L.
Harlocker, Susan L.
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Kalos, Michael
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59.5%;
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APPLICANT: Wang, Aljun
APPLICANT: Skelky, Yasir A.W.
APPLICANT: Hepler, William
ITITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
ITITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
ITITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
ITITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT PILLING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
SOFTWARE: PASTSEQ for Windows Version 3.0
SEQ ID NO 527
LENGTH: 320
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GENERAL INFORMATION:
APPLICANT: XL, Jiang
APPLICANT: Dillon, II
APPLICANT: Mitcham,
APPLICANT: Harlocker
APPLICANT: Harlocker
APPLICANT: Henderson
APPLICANT: Kalos, Mi
APPLICANT: Fanger, G
APPLICANT: Retter, W
APPLICANT: Stolk, Jc
APPLICANT: Day, Crai
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US-09-651-236-527
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; ORGANISM: Homo sapiens
US-09-759-143-527
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 Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
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                                                               Jiang, Yuqui
Henderson, Robert A.
Kalos, Michael D.
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Dillon, Davin C.
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Vedvick, Thomas
Carter, Darrick
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                                                                                                              Susan L.
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APPLICANT: Wang, Aijun
APPLICANT: Weaky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
ITITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
ITITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
ILLE REFERENCE: 210121.42718C18
CURRENT APPLICATION NUMBER: US/09/651,236
CURRENT FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 865
SOFTWARE: FRSTSEQ FOR Windows Version 3.0
SEQ ID NO 527
LENGTH: 320
TYPE: PAT
ORGANISM: Homo sapiens
US-09-651-236-527
; NUMBER OF SEQ ID NOS: 12
; SOPTWARE: PatentIn Ver. 2.
; SEQ ID NO 4
; LENGTH: 320
; TYPE: PRT
; ORGANISM: Homo Bapiens
US-09-968-033C-4
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                                                                                                                                      FILE REFERENCE: PF180P1
CURRENT APPLICATION NUMBER: US/09/968,033C
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: 60/237,275
PRIOR APPLICATION NUMBER: 60/237,275
PRIOR APPLICATION NUMBER: 09/339,115
PRIOR FILING DATE: 1999-06-24
PRIOR PILING DATE: 1999-06-24
PRIOR APPLICATION NUMBER: 09/053,303
PRIOR HILING DATE: 1998-04-01
PRIOR APPLICATION NUMBER: 08/465,980
PRIOR APPLICATION NUMBER: 08/465,980
PRIOR PILING DATE: 1995-06-06
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APPLICANT: Soppet et al.
TITLE OF INVENTION: Human Prostate Specific G-Protein Receptor HPRAJ70
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Best Local Similarity 59.5%; Pred. No. 2.1e-81;
Sest Local Similarity 55; Mismatches 67;
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Carter, Darrick
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                                                                                                                       ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-657-279-527
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SEQ ID NO 527
LENGTH: 320
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Matches
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Best Local
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APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C19
CURRENT PEPLICATION NUMBER: US/09/657,279
CURRENT FILING DATE: 2000-09-06
                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
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FastSEQ for Windows Version 3.0
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Mitcham, Jennifer L.
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                                                          60.4%; Score 979; DB 2; Length 320; Ilarity 59.5%; Pred. No. 2.1e-81; Conservative 55; Mismatches 67; Indels
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rick, Thomas
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US-10-012-896-527
; Sequence 527, Application US/10012896
; Patent No. 6943236
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                                                                                                                                                                                             FILE REPERENCE: 210121.427C27
CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 1011
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 527
LENGTH: 320
                                                                                                Query Match 60.4%; Score 979; DB 2; I Best Local Similarity 59.5%; Pred. No. 2.1e-81; Matches 182; Conservative 55; Mismatches 67;
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APPLICANT: Dil:
APPLICANT: Mit-
APPLICANT: Har:
APPLICANT: Jian
APPLICANT: Kal
                                                                                                                                                       TYPE: PRT
ORGANISM: Homo
-10-012-896-527
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
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Carter, Darrick
Li, Samuel X.
Aijun
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                                          RLFHVA 312
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Fanger, Gary R.
Wantanabe, Yoshihiro
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Mitcham, Jennifer L.
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Houghton, Raymond L.
Vinals de Bassols, Carlota
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Hepler, William T.
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Kalos, Michael D.
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US-09-680-728-3
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US-09-680-728-3
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SEQ ID NO 3
LENGTH: 32
TYPE: PRT
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Patent No. 6790631
GENERAL INFORMATION:
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Best Local (
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APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
FILE REFERENCE: 129.24USU1
CURRENT APPLICATION NUMBER: US/09/680,728
CURRENT FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR PILING DATE: 1999-10-05
NUMBER OF SEQ ID NOS: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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  304 AMFKIS
                             307 RLFHVA 312
                                                                                                               184
                                                                                                                                 188 MKLACDDIRVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSH 246
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                                                                                                                                                                                    128 AICHPLRHATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQDV 187
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                                                                                                                                                                                                                             64 MLAAIDLALSTSTMPKILALFWFDSRBITFDACLAQMFFIHALSAIBSTILLAMAFDRYV 123
                                                                                                                                                                                                                                                      68 MLSGIDILISTSSMEXMLAIFWENSTTIQFDACLLQIFAIHSLSGMESTVLLAMAFDRYV 127
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                                                         MKLAYTDTLPNVVYGLTAILLVMGVDVMFISLSYFLIIRAVLQLPSKSBRAKAFGTCVSH
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Daniel E.H. Afar
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    309
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                Sequence 2, Application US/09968033C
Patent No. 6824993
GENERAL INFORMATION:
APPLICANT: Soppet et al.
TITLE OF INVENTION: Human Prostate Specific G-Protein Receptor HPRAJ70
FILE REFERENCE: PF180P1
CURRENT APPLICATION NUMBER: US/09/968,033C
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: 60/237,275
PRIOR FILING DATE: 2000-10-03
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SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
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CURRENT APPLICATION NUMBER: US/10/017,066A
CURRENT FILING DATE: 2002-05-28
PRIOR APPLICATION NUMBER: US 09/680,728
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR PILING DATE: 1999-10-05
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APPLICANT: Steve Chappell Mitchell
APPLICANT: Douglas C. Saffran
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
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APPLICATION NUMBER: 09/339,115
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                                                                                                                                                                                                                                                                                                                                        AMPKIS 309
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59.5%;
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RESULT 17
US-08-465-980-2
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; LOCATION: (192)
; OTHER INFORMATION: Xaa equals any amino acid
US-09-968-033C-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2
Patent No.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve:
CURRENT APPLICATION DATA:
CURRENT APPLICATION UNHEER: US/08/465,980
FILING DATE: 06-JUN-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Perraro, Gregory D.
REGISTRATION NUMBER: 36,134
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PRIOR APPLICATION NUMBER: 09/053,303
PRIOR FILING DATE: 1998-04-01
PRIOR APPLICATION NUMBER: 08/465,980
PRIOR PILING DATE: 1995-06-06
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver: 2.1
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ORGANISM: Homo sapiens
FEATURE:
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APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: HUMAN G-PROTEIN RECEPTOR HPRAJ70
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                              CITY: Roseland
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                   STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       192 XPNVVYGLTAILLVMGVDVMFISLSYFLIIRTVLQLPSKSERAKAFGTCVSHIGVVLAFY 251
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o. 5756309
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Similarity 59.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VPLIGLSVVHRFGNSLHPIVRVVMGDIYLLLPPVINPIIYGAKTKQIRTRVLAMPKIS 309
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                                                                                                                                                                                                                                                                                                                     6 Becker Farm
                                                                                                                                                                                                                                                              USA
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STUART & OLSTEIN
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                                                                                                                                                                                                                                                                                                                                                            BAIN, GILFILLAN, CECCHI,
                                                                                                                                              Version #1.30
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TELECOMMUNICATION INFORMATION:

REFERENCE/DOCKET NUMBER: 325800-446

TELEPHONE:

201-994-1700

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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 320 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
PILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                         CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: HUMAN G-PROTEIN RECEPTOR HPRAJ70
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ELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                    STREET: 6 Beck
CITY: Roseland
STATE: New Jer
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TOPOLOGY: linear
            NAME: Ferraro, Gregory D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 32
                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI.
ADDRESSEE: STUART & OLSTEIN
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ilarity 59.4%;
Conservative 5
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Li, Yi
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                                                                                                               08/465,980
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                  325800-446
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Pred. No. 4.5e-80;
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RESULT 19
US-09-339-115-2
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Patent No. 6372891
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
ATTORNEY/AGENT INFORMATION:
NAME: Ferraro, Gregory D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 325800-446
TELECOMMUNICATION INFORMATION:
                                                                                                                        APPLICATION NUMBER: US/09/339,115
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
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TYPE: amino acid
TOPOLOGY: linear
                                                                                          APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                          CITY: Roseland
STATE: New Jersey
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Li, Yi
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Pred. No. 4.5e-80;
                                                                                                                                                                                                                      Version #1.30
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RESULT 20
US-09-680-728-4
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; TYPE: PRT
; ORGANISM: Homo Sapiens
US-09-680-728-4
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APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
APPLICANT: Mary Faris
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mitc
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                                                                                                                                                                                                               Query Match 59.5%; Score 964.5; DB 2; Best Local Similarity 59.4%; Pred. No. 4.5e-80; Matches 177; Conservative 54; Mismatches 66;
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Best Local Similarity
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SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: René S. Hubert
APPLICANT: Steve Chappell Mitchell
APPLICANT: Duglas C. Saffran
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR
TITLE OF INVENTION: UP-REGULATED IN PROSTATE CANCER AND USES THEREOF
FILE REFERENCE: 129.24USU1
CURRENT APPLICATION NUMBER: US/09/680,728
CURRENT FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/157,902
PRIOR FILING DATE: 1999-10-05
NUMBER OF SEQ ID NOS: 50
NUMBER OF SEQ ID NOS: 50
SOFTWARE: F8stSEQ for Windows Version 4.0
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SEQUENCE CHARACTERISTICS:
LENGTH: 320 amino acids
TYPE: amino acid
TYPE: amino acid
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  136 ATVLTLPRVTKIGVAAVVRGAALMAPLPVFIKQLPFCRSNILSHSYCLHQDVMKLACDDI 195
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                                                                                                                           72 LSTSTMPKILALFWFDSREISIEACLTQMFFIHALSAIESTILLAMAFDRYVAICHPLRH 131
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                                            LSTSTMPKILALFWFDSREISIEACLTQMFFIHALSAIESTILLAMAFDRYVAICHPLRH 131
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                                                                                                                                                                                                                 66; Indels
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196 RVNVVYGLIVIISAIGLDSLLISFSYLLILKTVLGL-TREAQAKAFGTCVSHVCAVFIFY 254
                                                                                                        132 AAVLNUTVTAQIGIVAVVRGSLFFFPLPLLIKRLAFCHSNVLSHSYCVHQDVMKLAYADT 191
                                                    192 LPNVVÝGLTAILLUMGUDUMFISLSYFLIIRTVLQLPSKSERAKAFGTCVSHIGVVLAFY 251
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